Page 1

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OM protein - protein search, using sw model

January 28, 2005, 16:54:57 Run on:

; Search time 40 Seconds (without alignments) 626.705 Million cell updates/sec

Title: Perfect score: Sequence:

US-10-048-071-28 1863 1 MIQPSINRTLFIHALNITKR......LTPGDBEESFIQLITPVRTN 378

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

478139 Total number of hits satisfying chosen parameters: 478139 seqs, 66318000 residues Searched:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

Issued Patents AA:*
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2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
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6: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		di			SUMMARIES		
Result No.	Score	Query Match	Length DB	80	. ДІ	Description	
7	1389	74.6	378	4	US-09-583-110-4405	i e	
7	863.5	46.3	384	4	US-09-107-532A-4556		
m	748.5	40.2	385	٣	US-09-134-001C-5166	Sequence 5166, Ap	
4	431		181	4	US-09-134-000C-5039		
5	350.5		368	4	US-09-252-991A-18904	189	
9	315	16.9	383	4	US-09-328-352-5969	Sequence 5969, Ap	
7	313		374	4	US-09-489-039A-10579		
80	307		375	4	US-09-543-681A-7400	Sequence 7400, Ap	
6	249.5		375	4	US-09-818-780-23		
10	232		303	4	US-09-198-452A-357		
11	131.5	7.1	141	4	US-09-134-000C-5040	5040,	
12	117.5	6.3	977	4	US-09-248-796A-15579		
13	115.5	6.2	800	m	US-08-776-265-3		
14	115.5	6.2	800	4	US-09-398-184-3	Sequence 3, Appli	
15	114.5	6.1	823	4	US-09-248-796A-16699		
16	113.5	6.1	922	4	US-09-883-134-9		
17	112.5	9	470	4	US-09-248-796A-23131		
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19	109.5	5.9	1753	4	US-09-248-796A-19154		
20	109	5.9	532	4	US-09-710-279-546	546,	
21	107.5	5.8	569	4	US-09-248-796A-16697		
22	107	5.7	563	ო	US-09-134-001C-3172	3172,	
23	106.5	5.7	1155	4	US-09-543-681A-6286	Sequence 6286, Ap	
24	104.5	5,6	1726	4	US-09-700-227-2		
25	103.5	5.6	564	4	US-09-107-532A-5248	524	
56	103.5	5.6	920	4	US-09-463-402-6		
27	103.5	5.6	921	4	US-09-889-572-4	4,	

Sequence 4971, Ap Sequence 404, App Sequence 402, App	5119 2303 10,	Sequence 10, Appl Sequence 10, Appl Sequence 10, Appl	-40	Sequence 15552, A Sequence 6, Appli Sequence 2586, Ap Sequence 4677, Ap
US-09-134-000C-4971 US-09-071-035-404 US-09-071-035-402	US-09-134-001C-5119 US-09-248-796A-23039 US-08-480-604A-10	US-08-915-136-10 US-08-957-310-10 US-10-011-366-10	US-09-084-517-10 US-09-538-092-326 US-09-543-681A-4674 US-09-134-000C-6159	US-09-248-796A-15552 US-08-856-253-6 US-09-710-279-2586 US-09-134-000C-4677
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1233 1416 1448	395 461 2366	2366 2366 2366	2366 960 385 666	380 512 611 910
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30 30 30	332		38 4 4 38 11 0 0 1	4 4 4 4 5 4 5 5 5 5 5 5 5 5 5 5 5 5 5 5

ALIGNMENTS

RESULT 1 US-09-583-110-4405

Sequence 4 Patent No. GENERAL IN APPLICANT TITLE OF TITLE	Sequence 4405, Application US/09583110 Patent No. 6699703 GENERAL INFORMATION: APPLICANT: Lynn Doucette-Stamm et al. TITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Relating to Streptococcus TITLE OF INVENTION: Nucleic Nucleic and Therapeutics CURRENT APPLICATION NUMBER: US 09/107,433 PRIOR PILING DATE: 1998-06-30 PRIOR FILING DATE: 1998-06-12 PRIOR FILING DATE: 1998-05-12 PRIOR FILING DATE: 1997-07-02 NUMBER OF SEQ ID NOS: 5322 IENGTH: 378 ITYPE: RAT TYPE: RA
Query Ma Best Loo Matches	Query Match Best Local Similarity 72.2%; Pred. No. 6.7e-125; Matches 273; Conservative 50; Mismatches 55; Indels 0; Gaps 0;
o o	1 MIQFSINRTLFIHALNTTKRAISTKNÄIPILSSIKIEVTSTGVTLTGSNGOISIENTIPV 60
& 8	61 SNENAGLLITSPGAILLEASFFINISSLPDISINVKEIBOHOVVLTSGKSEITLKGKDV 120
දු පු	121 DOYPRLOBUSTENPLILKTKALKSIIABTAFAASLOBSRPILTGVHIVLSNHKDFKAVAT 180
දු පු	181 DSHRMSQRLITLDNTSADLAVVLPSKSRRRFSAVFTDDIBTVEVPFSPSQILFRSEHISF 240
දි සි	241 YTRLLEGNYPDTDRLLMTEFETEVVFNTQSLRHAMERAFLISNATONGTVKLEITQNHIS 300
È	301 AHVNSPEVGKVNEDLDIVSQSGSDLTISPNPTYLIESLKAIKSETVKIHFLSPVRPPTLT 360

```
GENERAL INFORMATION:

APPLICANT: LyTH DOUGE LEGISTARM Et al

TITLE OF INVENTION: NUCLBIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUE

TITLE OF INVENTION: BEIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: GTC-007

CURRENT APPLICATION NUMBER: US 60/064,964

PRIOR PILING DATE: 1999-08-13

PRIOR PELICATION NUMBER: US 60/064,964

PRIOR FILING DATE: 1997-11-08

PRIOR FILING DATE: 1997-11-08

RIOR FILING DATE: 1997-08-14

NUMBER OF SEQ ID NOS: S674

SEQ ID NO 5166

FEATURE OF SEG ID NOS: S674
                                                        241 YTRLLEGNYPDTDRLLMTEFETEVVFNTQSLRHAMERAFLISNATQNGTVTLEITQNHIS 300
                                                                                                                                                                         247 YSRLLEGNYPDTNRLIPSSFNTEVEFSVPSFLAAIBRASLLSHEGRNNIVRLSIRPDAVV 306
                                                                                                                                                                                                                          301 AHVNSPEVGKVNEDLDIVSQSGSDLTISFNPTYLIESLKAIKSETVKIHFLSPVRPFTLT 360
                                                                                                                                                                                                                                                   61 SNENAGLL-ITSPGAILLEASFFINIISSLPDISINVKEIEQHQVVLTSGKSBITLKGKD 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            69 QVDGBEIVEITETGSVVLPGRPPVDIIKKLPGKEVKLSTNEQPQTLITSGHSEFNLSGLD 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            120 VDQYPRLQEVSTENPLILKTKLLKSIIAETARAASLQESRPILTGVHIVLSNHKDPKAVA 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             129 PDQYPLLPEVSRDDAIQLSVKVLKNIIAQTNPAVSTSETRPVLTGVNWLIQDN-ELICTA 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                180 IDSHRMSQRLITLDNTSADLMVVLPSKSLREFSAVFTDDIETVEVFFSFSQILFRSEHIS 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TDSHRLAVRKIQLEDESENKAVIIPGKALSELNKIMSDSDEDIDIFFASNQVLFRVGNIN 247
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              248 PISRLLEGHYPDTTRLPPENYEIKLGINNGDPYHAIDRASLLAREGGNNVIKLSTGNELV 307
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                300 SAHVNSPEVGKVNEDLDIVSQSGSDLTISFNPTYLIESLKAIKSETVKIHFLSPVRPFTL 359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DSHRMSQRLITLDNTSADLMVVLPSKSLREFSAVFTDDIETVEVFFSPSQILFRSEHISF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 MIQPSINRTLFIHALNTTKRAISTKNAIPILSSIKIBVTSTGVTLTGSNGQISIENTIPV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          40.2%; Score 748.5; DB 3; Length 385; 40.5%; Pred. No. 2.4e-63; Live 86; Mismatches 136; Indels 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 5166, Application US/09134001C Patent No. 6380370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Staphylococcus epidermidis
                                                                                                                                                                                                                                                                                                                           361 PGDEEESFIQLITPVRTN 378
                                                                                                                                                                                                                                                                                                                                                            367 PTEDGVQFIQLITPVRTN 384
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 3
US-09-134-001C-5166
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                                                                                                                                                                                                                                                                                       APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
301 AHVHSPEVGKVNEEIDTDQVTGEDLTISFNPTYLIDSLKALNSEKVTISFISAVRPFTLV 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 SNENAGLLITSPGAILLEASFFINIISSLPDISINVKEIEQHQVVLTSGKSEITLKGKDV 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             YPRIOEVSTENPLILKTKLLKSIIAETAFASLOESRPILITGVHIVLSNHKDFKAVAT 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 46.3%; Score 863.5; DB 4; Length 384; Best Local Similarity 46.6%; Pred. No. 2.1e-74; Matches 176; Conservative 72; Mismatches 129; Indels 1.
                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
STREET: 100 Beaver Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/107,532A
PILING DATE: 30-Jun-1998
PRIOR APPLICATION NUMBER: 60/085,598
PILING DATE: 14 May 1998
APPLICATION NUMBER: 60/051571
RILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc feature
LOCATION: (B) LOCATION 1...384
SEQUENCE DESCRIPTION: SEQ ID NO: 4556:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Enterococcus faecium
                                                                                                                                                                                                                    Sequence 4556, Application US/09107532A
Patent No. 6583275
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: CD/ROM ISO9660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 384 amino acids
                                                   361 PGDEESFIQLITPVRTN 378
                                                                            361 PADIDEDFMOLITPVRIN 378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 4556:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (781)893-8277
                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 7310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: protein
HYPOTHETICAL: YES
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Waltham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 02354
                                                                                                                                                                                             US-09-107-532A-4556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-107-532A-4556
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Total number of hits satisfying chosen parameters: 824507 seqs, 355394441 residues Searched:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

1seued Patents NA:*
1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
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5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:* Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

			de			SUMMARIES	
Result No.	ult No.	Score	Query Match	Length DB	DB	ID	Description
	н	849	74.9	3200	۳	US-09-381-862-2	
	N	608.2	53.6	1137	4	122	
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	4	317	28.0	1155	٠ 4	TIS-08-301-32/-20	20,
	Ŋ	247.4	21.8	1158	۰,	TIS-09-134-001	50
	v	236.2	20.8	2347	4	US-03-134-001C-2329	2329,
	7	236.2	20.8	2347	4	TIS-08-701-0068 153	153,
	œ	159.6	14.1	546	4	US-09-134-0007-153	
O	σ	59.2	5.2	640681	4	US-09-790-988-1	1634,
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υ	11	54.8	8.4	1141	٠ ٦	TIS-09-806-1008-22	22
	12	49.6	4	426	٠ ٦	•	
	13	40	. 4	107077	۲ -	US-03-134-000C-1635	19
t	14	2 - 4		140041	4 .	US-09-790-988-1	H
,	۲ u	*	4. 4	14066	4	US-09-601-198-56	Sequence 56, Appl
) (7.0	7	471	4.	US-09-543-681A-1232	5
	9 1	45.8	4.0	751	4	US-08-956-171E-892	1 0
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	18	45.6	4.0	832	4	US-09-621-976-2813	
	19	45.4	4.0	821	ო	US-08-998-416-541	•
	20	45	4.0	1134	4	US-09-601-198-62	1 0
	21	44.8	4.0	423	4	US-09-710-279-1771	
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	23	44.8	. 4	1 5	• •	110 00 10 1 1 1 45	
	70	0 0 0		# L	ი •	US-US-134-001C-2705	Seguence 2705, Ap
	* u			3315	4.	US-09-710-279-3820	
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	Sequence 82, Appl	Semionce as	יייייייייייייייייייייייייייייייייייייי	pednemce T' Appli	Segmence 1 April:	Tradit / Commission	addresse to Appl	Sequence 208, App	The Cold of the Cold		Segmence 5237 An	, .	peduence I, Appli	Segmence 11532 A	•	sednence 444, App	Segmence 23, April	TATE COLORDON	ddw '/or apmanha	Segmence 3017. An	C COCCOMON	acqueires s, Appli	Sequence 6. Appli	C	יישרתיים ליי שלחלת	Sequence 1, Appli
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ALIGNMENTS

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                                   Length 3200;
                                                    10; Indels
                               Score 849; DB 3; L
Pred. No. 9.5e-204;
0; Mismatches 10;
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 Clinical Isolate SP-7-44
                               Query Match
Best Local Similarity 98.7%;
Matches 866; Conservative
; STRAIN:
US-09-381-862-2
                            Query Match
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APPLICANT: Lynn Doucette-Stamm et al.

TITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Relating to Streptococ FILE OF INVENTION: Pneumoniae for Diagnostics and Therapeutics FILE REFERENCE: PATHOL-07A
CURRENT FILING DATE: 2006-05-26
PRIOR PILING DATE: 1908-06-30
PRIOR PILING DATE: 1908-06-30
PRIOR PILING DATE: 1998-06-30
PRIOR PILING DATE: 1998-06-30
PRIOR FILING DATE: 1998-06-30
PRIOR FILING DATE: 1998-05-12
PRIOR FILING DATE: 1997-07-02
NUMBER OF SEQ ID NOS: 5322
SEQ ID NO 1744
LENGTH: 1137
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Pred. No. 2e-143;
0; Mismatches 328; Indels 0;
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Best Local Similarity 71.1%;
Matches 805; Conservative
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Sequence 1744, Application US/09583110

RESULT 2 US-09-583-110-1744

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7676 GICGIAATICCTAGCCGITCICIACGCGAATITICAGCGGTATITACAGAIGATATACGAA 7735
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Patent No. 6420135
GENERAL INFORMATION:
APPLICANT: Charles Kunsch
TITLE OF INVENTION: Streptococcus pneumoniae Polynucleotides and Sequences
NUMBER OF SEQUENCES: 391
CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
STATE: Maryland
                                                                                                                                                                                                                                 1020
                                                                                                                                                                                                  961 TCTGGTAGTGATTTAACTATCAGCTTCAACCTAACTTACCTTATTGAGTCTTTAAAAGCT 1020
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                                                                                                     901 GCCCATGTTCACTCTCCAGAAGTTGGTAAAGTAAACGAAAATCGATACTGATCAGGTT 960
                                               781 AACACTACTATTTATTTAATGTGGTAAACTTTACGCCAGTCAATGGAGCGTGCCCGTCTT 840
                                                                                   841 ATTICTAAIGCTACTCAAAAIGGTACTGTTAAGCTTGAGATTACTCAAAATCATATTTCA 900
                                                                                                                                            GCTCATGTTAACTCACCTGAGGTTGGTAAGGTAAACGAGGATTTAGATATTGTTAGTCAG 960
721 TATACTCGTCTCCTAGAAGGAAACTATCCTGATACAGATCGCTTGATTCCAACAGACTTT 780
                             GAGACGGAGGTTGTTTTCAATACCCAATCCCTTCGCCACGCTATGGAACGTGCCTTCTTG
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MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/961,527
FILING DATE:
CLASSIFICCATION DATA:
APPLICATION NUMBER: BAPLICATION NUMBER: FILING DATE:
FILING DATE:
FILING DATE:
TELEBRING PART:
TELEBRING NOMBER: 36,33
REGISTRATION NUMBER: 36,33
REGISTRATION NUMBER: 36,349
REGISTRATION NUMBER: 36,349
REGISTRATION NUMBER: 36,349
TELEBRINGE: (301) 309-8504
TELEBRINGE: (301) 309-8512
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 53.6%; Score 608.2; DB 4; Best Local Similarity 71.1%; Pred. No. 5.7e-143; Matches 805; Conservative 0; Mismatches 328;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 21338 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
TOPOLOGY:
US-08-961-527-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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us-10-048-071-27.rni 4 10:26:48 2005 Fe.

7615 7136 GCTATTAGTTCTAAAAATGCCATTCCTATTTTATCAACAGTAAAAATTGACGTGACCAAT 7195 7256 AAAAATGAAGATGCTGGTTTGTTAATTACTTCTTTAGGTTCGATCCTTCTTGAAGCTTCT 7315 7316 ircirirarcaargractarcrastriaccrsarsraacrcrissrrraarsaastasa 7375 7496 TTACTCAAGAAATTATTAATGAAACAGCCTTTGCTGCAAGTACACAAGAGAGTCGTCGC 7555 7196 gaadgiaitracitiraltiggicaaatggicaariticaatigaaaattiratitracicaa 7255 421 TIATTGAAGTCTATTATTGCTGAAACAGCTTTTGCAGCCAGTTTACAAGAAAGTCGTCCT 480 481 ATTTTAACAGGAGTTCATATTGTATTAAGTAATCATAAAGATTTTAAAGCAGTAGCGACT 540 7436 GAACAATATCCACGAATCCAAGAAATTTCAGCAAGCACTCCTTTAATACTTGAAACAAAA CAACACCAAGTTGTTTTAACCAGTGGTAAATCAGAGATTACCTTTAAAAGGAAAAGATGTT 181 AGTAATGAAAATGCTGGTTTGCTAATTACCTCTCCAGGAGCTATTTTATTAGAAGCTAGT TITITIATIAATATITICAAGTITGCCAGATATTAGTATAAAATGTTAAAGAAATTGAA GACCAGTATCCTCGTCTACAAGAAGTATCAACAGAAAATCCTTTGATTTTAAAAACAAAA 7076 ATGATTCATTTTCAATTAATAAAATTTATTTCTACAAGCATTAAATACTACTAAGAGA 61 GCTATTAGCACTAAAAATGCCATTCCTATTCTTTCATCAAAAAATTGAAGTCACTTCT 121 ACAGGAGTAACTTTAACAGGGTCTAAACGGTCAAATATCAATTGAAAACACTATTCCTGTA 7556 ATTITIAACAGGGGCCACTTCGTATTGAGTCAACACAAAGAGGTTAAAAACAGTTGCAACA 7616 GACTCTCATCGCCTAAGCCAGAAAAAATTGACTCTTGAAAAAAATAGTGATGATTTTGAT 1. ATGATTCAATTTTCAATTAATCGCACATTATTTATTCATGCTTTAAATACAACTAAACGT 541 GACTCTCATCGTATGAGCCAACGTTTAATCACTTTGGACAATACTTCAGCAGATTTGATG

7855 7856 AACACTACTATTACTTTTAATGTGGTAAACTTACGCCAGTCAATGGAGCGTGCCCGCCTT 7915 TTATCAAGTGCGACTCAAAATGGTACTGTGAAACTTGAAATTAAGGATGGGGTTGTTAGC 7975 7736 ACTGTAGAGATTTTCTTTGCCAATAACCAAATCCTCTTTAGAAGCGAAAATATTAGCTTC 7795 1021 ATTAAAAGTGAAACAGTAAAAATTCATTTCTTATCACCAGTTCGACCATTCACCCTAACA 1080 840 3096 Traaaragccaaaagcrcacrarraccritarcrcaccrctrccrccaritacrcrrcrg 8155 780 841 ATTICIBATGCTACTCAAAATGGTACTGTTAAGCTTGAGATTACTCAAAATCATATTCA 900 GCTCATGTTAACTCACCTGAGGTTGGTAAGGTAAACGAGGATTTAAGATATTGTTAGTCAG 960 7796 TATACTCGTCTCCTAGAAGGAAACTATCCTGATACAGATCGCTTGATTCCAAGAGAGACTTT 781 GAGACGGAGGTTGTTTCAATACCCAATCCCTTCGCCACGCTATGGAACGTGCCTTCTTG TCTGGTAGTGATTTAACTATCAGCTTCAATCCAACTTACCTTATTGAGTCTTTAAAAGCT 661 ACCGTTGAGGTATTTTTCTCACCAAGCCAAATCTTGTTCAGAAGTGAACACACATTTCTTTT TATACACGCCTCTTAGAAGGAAATTATCCCGATACAGACCGTTTATTAATGACAGAATTT

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(FILE 'CAPLUS' ENTERED AT 09:49:53 ON 04 FEB 2005)
L1
            455 S (DNAN OR DNA N)
L2
              2 S L1 AND (STREPTOCOCC? OR S) (W) PYOGEN?
     (FILE 'REGISTRY' ENTERED AT 09:52:36 ON 04 FEB 2005)
              1 SEA FILE=REGISTRY ABB=ON PLU=ON "DNA (STREPTOCOCCUS PYOGENES
L4
                GENE DNAN)"/CN
     FILE 'CAPLUS' ENTERED AT 09:54:56 ON 04 FEB 2005
L5
              1 S L4
              2 S L2 OR L5
L6
     ANSWER 1 OF 2 CAPLUS COPYRIGHT 2005 ACS on STN
L6
     Entered STN: 03 May 2002
ACCESSION NUMBER:
                         2002:332364 CAPLUS
                         136:352018
DOCUMENT NUMBER:
TITLE:
                         Streptococcus pyogenes DNA
                         polymerase III holoenzyme subunits and their genes
INVENTOR(S):
                         McHenry, Charles S.; Bullard, James M.; Janjic,
                         Nebojsa; Manhardt, Erika L.; Kery, Vladimir; Williams,
                         Jennifer C.
                         Replidyne, Inc., USA
PATENT ASSIGNEE(S):
SOURCE:
                         PCT Int. Appl., 268 pp.
                         CODEN: PIXXD2
DOCUMENT TYPE:
                         Patent
                         English
LANGUAGE:
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
     PATENT NO.
                         KIND
                                DATE
                                            APPLICATION NO.
                                                                    DATE
                         ----
                                _____
                                            _____
     WO 2002034936
                         A2
                                20020502
                                            WO 2001-US48396
                                                                    20011029
     WO 2002034936
                          C2
                                20030417
     WO 2002034936
                         Α3
                                20020725
         W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN,
           CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
             GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR,
             LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT,
             RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US,
             UZ, VN, YU, ZA, ZW
         RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AM, AZ, BY, KG,
             KZ, MD, RU, TJ, TM, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR,
             IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN,
             GQ, GW, ML, MR, NE, SN, TD, TG
     AU 2002032586
                          Α5
                                20020506
                                            AU 2002-32586
                                                                    20011029
PRIORITY APPLN. INFO .:
                                            US 2000-244023P
                                                                    20001027
                                            WO 2001-US48396
                                                                 W 20011029
     Streptococcus pyogenes nucleic acid mols. encoding
AB
     polC, dnaE (subunit \alpha), holA (subunit \delta), holB (subunit
     \delta'), dnaX (subunit \tau), dnaN (subunit \beta), SSB
     (single-stranded DNA-binding protein), dnaG (primase), dnaQ (subunit
     \epsilon), dnaA and dnaB proteins, as well as nucleic acid mols.
     comprising the oriC origin of replication are provided. The encoded
     subunit proteins of S. pyogenes DNA polymerase III are
     also provided. The nucleic acid mols. and proteins are useful for
     reconstituting replicases and polymerases for sequencing, amplification,
```

and screening for compds. which modulate the function of the polymerase or replicase.

IT 419927-96-1, DNA (Streptococcus pyogenes gene dnaN)

RL: ANT (Analyte); BSU (Biological study, unclassified); BUU (Biological use, unclassified); PRP (Properties); THU (Therapeutic use); ANST (Analytical study); BIOL (Biological study); USES (Uses) (nucleotide sequence; Streptococcus pyogenes DNA polymerase III holoenzyme subunits and their genes)

L6 ANSWER 2 OF 2 CAPLUS COPYRIGHT 2005 ACS on STN

ED Entered STN: 09 Feb 2001

ACCESSION NUMBER: 2001:101168 CAPLUS

DOCUMENT NUMBER: 134:143863

TITLE: DNA replication proteins of Gram-positive bacteria and

their use to screen for chemical inhibitors

INVENTOR(S): O'donnell, Michael E.; Bruck, Irina; Zhang, Dan;

Whipple, Richard

PATENT ASSIGNEE(S): The Rockefeller University, USA

SOURCE: PCT Int. Appl., 238 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 2

PATENT INFORMATION:

PATENT NO.	KIND DATE	APPLICATION NO.	DATE
WO 2001009164	A2 200102	08 WO 2000-US20666	20000728
		A, BB, BG, BR, BY, CA, I, GB, GD, GE, GH, GM,	
•	· · · · · · · · · · · · · · · · · · ·	R, KZ, LC, LK, LR, LS,	
MD, MG, MK SK, SL, TJ	• • • • • •	O, NZ, PL, PT, RO, RU,	SD, SE, SG, SI,
		D, SL, SZ, TZ, UG, ZW,	AT, BE, CH, CY,
· · · · · · · · · · · · · · · · · · ·		R, IE, IT, LU, MC, NL, W, ML, MR, NE, SN, TD,	
• •	•	19 AU 2000-67499	20000728
US 2003129633 PRIORITY APPLN. INFO.:	A1 200307	10 US 2002-282287 US 1999-235245	
PRIORITI AFFUN. INFO		US 1999-146178P	P 19990729
		US 1998-74522P US 1998-93727P	
		WO 2000-US20666	

AB The present invention relates to α -large, α -small, δ , δ ', τ , β , SSB, DnaG, and DnaB and (polC, dnaE, holA, holB, dnaX, dnaN, ssb, dnaG, dnaB) genes encoding them from Gram pos. bacteria, preferably Streptococcus pyogenes and Staphylococcus aureus. The individual genes and proteins or polypeptides are useful in identification of compds. With antibiotic activity. Thus, the structure and mechanism of the chromosomal replicase of S. pyogenes and S. aureus have been elucidated. These DNA polymerases use a sliding clamp (the dnaN-encoded β subunit) and clamp loader (the dnaN-encoded τ subunit). The clamp and clamp loader components of Gram-neg. cells could be exchanged for those of Gram-pos. cells.

(FILE 'MEDLINE, BIOSIS, EMBASE, WPIDS, CONFSCI, SCISEARCH, JICST-EPLUS, JAPIO' ENTERED AT 09:56:11 ON 04 FEB 2005)

L7 1 S L6

L7 ANSWER 1 OF 1 WPIDS COPYRIGHT 2005 THE THOMSON CORP on STN

ACCESSION NUMBER: 2001-147453 [15]

C2001-043718

DOC. NO. CPI: TITLE:

Isolated DNA molecule from a Gram positive bacterium encoding DNA replication proteins used to identify

compounds which have antibiotic activity.

WPIDS

DERWENT CLASS:

B04 D16

INVENTOR(S):

BRUCK, I; O'DONNELL, M E; WHIPPLE, R; ZHANG, D

PATENT ASSIGNEE(S):

(UYRQ) UNIV ROCKEFELLER

COUNTRY COUNT:

91

PATENT INFORMATION:

PATENT NO KIND DATE WEEK LA PG

WO 2001009164 A2 20010208 (200115)* EN 238

RW: AT BE CH CY DE DK EA ES FI FR GB GH GM GR IE IT KE LS LU MC MW MZ

NL OA PT SD SE SL SZ TZ UG ZW

W: AE AL AM AT AU AZ BA BB BG BR BY CA CH CN CR CU CZ DE DK DM EE ES FI GB GD GE GH GM HR HU ID IL IN IS JP KE KG KP KR KZ LC LK LR LS

LT LU LV MA MD MG MK MN MW MX NO NZ PL PT RO RU SD SE SG SI SK SL

TJ TM TR TT TZ UA UG US UZ VN YU ZA ZW

AU 2000067499 A 20010219 (200129)

APPLICATION DETAILS:

PATENT NO	KIND	APPLICATION	DATE
WO 2001009164	A2	WO 2000-US20666	20000728
AU 2000067499	A	AU 2000-67499	20000728

FILING DETAILS:

PATENT NO	KI	ND	PATENT NO
	·		
AU 2000067499	Α	Based on	WO 2001009164

PRIORITY APPLN. INFO: US 1999-146178P 19990729

AN 2001-147453 [15] WPIDS

AB WO 200109164 A UPAB: 20011129

NOVELTY - Isolated DNA molecule (I) from a Gram positive bacterium comprises a coding region from a polC, dnaE, holA, holB, dnaX, dnaN, ssb, dnaG or a dnaB gene.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are also included for the following:

- (1) an expression system comprising an expression vector into which is inserted (I);
 - (2) a host cell comprising (I);
- (3) an isolated protein or polypeptide from a Gram positive bacterium which is alpha-large, alpha-small, delta, delta prime, tau, beta, SSB, DnaG or DnaB protein or polypeptide; and
 - (4) a method of identifying compounds which inhibit the activity of a

polymerase product of polC or dnaE comprising:

- (a) forming a reaction mixture containing a primed DNA molecule, a polymerase product of polC or dnaE, a candidate compound, a dNTP, and optionally a beta subunit and/or a tau complex, where at least one of the polymerase product of polC or dnaE, beta subunit, tau complex, or a subunit or combination of subunits is derived from a Eubacteria other than Escherichia coli;
- (b) subjecting the reaction mixture to conditions effective for polymerization extension products in the absence of the candidate compound;
- (c) analyzing the reaction mixture for the presence or absence of nucleic acid polymerization extension products; and
- (d) identifying the candidate compound in the reaction mixture where there is an absence of nucleic acid polymerization products.

USE - (I) encodes proteins that replicate the chromosome of Gram positive bacteria and are used for sequencing and amplification of DNA and in drug discovery to identify compounds which have antibiotic activity through interference with replication. The methods identify compounds that are active at the level of DNA replication and result in arrest of cell growth or cell death of bacteria to treat bacterial infections in animals.

ADVANTAGE - (I) encodes proteins which provide further targets for antibiotics. The methods are amenable to high throughput screening assays. Dwg.0/21

(FILE 'USPATFULL' ENTERED AT 09:57:14 ON 04 FEB 2005)

19 S L5 OR L1(L) ((STREPTOCOCC? OR S) (W) PYOGEN?)

L9 ANSWER 1 OF 19 USPATFULL on STN

L9

ACCESSION NUMBER: 2005:30758 USPATFULL

TITLE: Microbial operons

INVENTOR(S): Wang, Liangsu, San Diego, CA, UNITED STATES
Zamudio, Carlos, La Jolla, CA, UNITED STATES

NUMBER DATE

PRIORITY INFORMATION: US 2003-474768P 20030529 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: KNOBBE MARTENS OLSON & BEAR LLP, 2040 MAIN STREET,

FOURTEENTH FLOOR, IRVINE, CA, 92614

NUMBER OF CLAIMS: 35 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 11 Drawing Page(s)

LINE COUNT: 10876

AB Described herein is a method for predicting operons in prokaryotes. Also described herein are vectors comprising operons predicted using the this method as well as methods of using antisense nucleic acids complementary to at least a portion of a predicted proliferation-required operon to inhibit cellular proliferation. Methods of using such antisense nucleic acids to sensitize cells for use in assays to identify compounds which possess the ability to inhibit cellular proliferation are also

described.

INCLM: 435/006.000 INCL

INCLS: 702/020.000

NCLM: 435/006.000 NCL

NCLS: 702/020.000

ANSWER 2 OF 19 USPATFULL on STN

ACCESSION NUMBER: 2005:10919 USPATFULL

Fragmentation-based methods and systems for de novo TITLE:

sequencing

Boecker, Sebastian, Bielefeld, GERMANY, FEDERAL INVENTOR(S):

REPUBLIC OF

Boom, Dirk van den, La Jolla, CA, UNITED STATES

KIND NUMBER DATE US 2005009053 A1 20050113

PATENT INFORMATION:

US 2004-830943 A1 20040422 (10) APPLICATION INFO.:

NUMBER DATE

PRIORITY INFORMATION: US 2003-466006P 20030425 (60)

DOCUMENT TYPE: Utility APPLICATION FILE SEGMENT:

Stephanie L. Seidman, FISH & RICHARDSON P.C., 12390 El LEGAL REPRESENTATIVE:

Camino Real, San Diego, CA, 92130-2081

NUMBER OF CLAIMS: 84 EXEMPLARY CLAIM: 1

17 Drawing Page(s) NUMBER OF DRAWINGS:

LINE COUNT: 4217

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods and systems, particularly mass spectrometric methods and

systems, for the analysis and sequencing of biomolecules, particularly

nucleic acids, by fragmentation are provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCLM: 435/006.000 INCL

INCLS: 435/007.100; 702/020.000

NCL NCLM: 435/006.000

NCLS: 435/007.100; 702/020.000

ANSWER 3 OF 19 USPATFULL on STN

2004:299897 USPATFULL ACCESSION NUMBER:

System for discovery of agents that block yersinia TITLE:

pestis and pseudomonas aeruginosa dna replication

INVENTOR(S): Bullard, James M., Longmont, CO, UNITED STATES

Janjic, Nebojsa, Boulder, CO, UNITED STATES

McHenry, Charles S., Denver, CO, UNITED STATES KIND DATE NUMBER

US 2004235766 A1 US 2003-476597 A1 WO 2002-US15111 20041125 PATENT INFORMATION: APPLICATION INFO.: 20031031 (10)20020514

> Shears 571-272-2528 Searcher :

NUMBER DATE PRIORITY INFORMATION: US 2001-60290725 20010514 US 2001-60332644 20011105 DOCUMENT TYPE: Utility FILE SEGMENT: APPLICATION SWANSON & BRATSCHUN L.L.C., 1745 SHEA CENTER DRIVE, LEGAL REPRESENTATIVE: SUITE 330, HIGHLANDS RANCH, CO, 80129 NUMBER OF CLAIMS: EXEMPLARY CLAIM: 77 Drawing Page(s) NUMBER OF DRAWINGS: LINE COUNT: 5150 CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Y. pestis and P. aeruginosa nucleic acid molecules encoding dnaE, holA, holB, holC, holD, holE, dnaX, dnaN, SSB, dnaG, dnaQ, proteins are provided. The encoded proteins are also provided. The nucleic acid molecules and proteins are useful for reconstituting replicases and polymerases for sequencing, amplification, and screening for compounds

which modulate the function of the polyemersase or replicase.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCLM: 514/044.000 INCL INCLS: 435/006.000 NCL NCLM: 514/044.000 NCLS: 435/006.000

ANSWER 4 OF 19 USPATFULL on STN

2004:250212 USPATFULL ACCESSION NUMBER:

Nucleic acid and amino acid sequences relating to TITLE:

Streptococcus pneumoniae for diagnostics and

therapeutics

Doucette-Stamm, Lynn A., Framingham, MA, United States Bush, David, Somerville, MA, United States INVENTOR(S):

Genome Therapeutics Corporation, Waltham, MA, United PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE ______ US 6800744 PATENT INFORMATION: B1 20041005 US 1998-107433 19980630 (9) APPLICATION INFO.:

NUMBER DATE ______ US 1998-85131P 19980512 (60) US 1997-51553P 19970702 (60) PRIORITY INFORMATION:

DOCUMENT TYPE: Utility FILE SEGMENT: GRANTED

PRIMARY EXAMINER: Brusca, John S. Zhou, Shubo "Joe " ASSISTANT EXAMINER:

LEGAL REPRESENTATIVE: Genome Therapeutics Corporation

NUMBER OF CLAIMS: 14 EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 0 Drawing Figure(s); 0 Drawing Page(s)

LINE COUNT: 11545

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The invention provides isolated polypeptide and nucleic acid sequences

571-272-2528 Searcher : Shears

derived from Streptococcus pneumonia that are useful in diagnosis and therapy of pathological conditions; antibodies against the polypeptides; and methods for the production of the polypeptides. The invention also provides methods for the detection, prevention and treatment of pathological conditions resulting from bacterial infection.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCL INCLM: 536/023.100

INCLS: 435/006.000; 435/320.100; 435/325.000; 435/254.000; 435/419.000;

536/024.100; 536/023.400; 536/024.320

NCL NCLM: 536/023.100

NCLS: 435/006.000; 435/320.100; 435/325.000; 435/419.000; 536/023.400;

536/024.100; 536/024.320

L9 ANSWER 5 OF 19 USPATFULL on STN

ACCESSION NUMBER: 2004:217811 USPATFULL

TITLE: Development of novel anti-microbial agents based on

bacteriophage genomics

INVENTOR(S): Pelletier, Jerry, 8 Lakeview, Baie-D'Urfe, Quebec,

CANADA H9X 3B1

Gros, Philippe, 107 Montrose, St. Lambert, Quebec,

CANADA J4R 1X4

DuBow, Michael, 4901 Coolbrook Avenue, Montreal,

Quebec, CANADA H3X 2K8

NUMBER KIND DATE

PATENT INFORMATION: US 6783930 B1 20040831 APPLICATION INFO.: US 1999-454252 19991202 (9)

RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 1999-407804, filed

on 28 Sep 1999

NUMBER DATE

PRIORITY INFORMATION: US 1998-110992P 19981203 (60)

DOCUMENT TYPE: FILE SEGMENT: Utility GRANTED

PRIMARY EXAMINER:
ASSISTANT EXAMINER:

Wax, Robert A. Mitra, Rita

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 10 1

NUMBER OF DRAWINGS:

9 Drawing Figure(s); 8 Drawing Page(s)

LINE COUNT: 9158

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB A method for identifying suitable targets for antibacterial agents based on identifying targets of bacteriophage-encoded proteins is described. Also described are compositions useful in the identification methods and in inhibiting bacterial growth, and methods for preparing and using such compositions.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCL INCLM: 435/005.000

INCLS: 435/007.100; 435/007.330; 435/007.800; 435/883.000; 536/023.700;

530/350.000; 530/820.000

NCL NCLM: 435/005.000

NCLS: 435/007.100; 435/007.330; 435/007.800; 435/883.000; 530/350.000;

530/820.000; 536/023.700

ANSWER 6 OF 19 USPATFULL on STN

2004:178350 USPATFULL ACCESSION NUMBER:

TITLE: DNA sequences from staphylococcus aureus bacteriophage

44AHJD that encode anti-microbial polypeptides

Pelletier, Jerry, Baie-D'Urfe, CANADA INVENTOR(S):

Gros, Philippe, St. Lambert, CANADA

DuBow, Michael, Antony, FRANCE

Bergeron, Dominique, Montreal, CANADA

Phagetech, Inc. (non-U.S. corporation) PATENT ASSIGNEE(S):

> NUMBER DATE KIND ----- -----

US 2004137516 A1 20040715 US 2003-449830 A1 20030531 (10) PATENT INFORMATION:

APPLICATION INFO.:

Continuation-in-part of Ser. No. US 2000-727892, filed RELATED APPLN. INFO.:

on 1 Dec 2000, PENDING Continuation of Ser. No. WO

2001-CA1754, filed on 30 Nov 2001, UNKNOWN

NUMBER DATE

PRIORITY INFORMATION: US 1999-168777P 19991201 (60)

DOCUMENT TYPE: Utility APPLICATION FILE SEGMENT:

SUGHRUE MION, PLLC, 2100 PENNSYLVANIA AVENUE, N.W., LEGAL REPRESENTATIVE:

SUITE 800, WASHINGTON, DC, 20037

43 NUMBER OF CLAIMS: 1 EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 24 Drawing Page(s)

LINE COUNT: 3666

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

This invention relates to newly identified polynucleotides and polypeptides, and their production and uses, as well as their variants, agonists and antagonists, and their uses. In particular, the invention relates to specific interaction between the S. aureus STAAU R2 related protein or specific regions thereof, and growth-inhibitory proteins encoded by the S. aureus bacteriophage genome. The invention relates to the use of these interaction target sites as the basis of drug screening assays.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCL INCLM: 435/007.100 NCLM: 435/007.100 NCL

L9 ANSWER 7 OF 19 USPATFULL on STN

ACCESSION NUMBER: 2004:152148 USPATFULL

Retroductal salivary gland genetic vaccination TITLE: Tucker, Sean, San Francisco, CA, UNITED STATES INVENTOR(S):

Bennett, Michael, El Sobrante, CA, UNITED STATES

Chen, Yen-Ju, Alameda, CA, UNITED STATES Olson, David, Alameda, CA, UNITED STATES

Genteric, Inc., Alameda, CA (U.S. corporation) PATENT ASSIGNEE(S):

> NUMBER KIND DATE ______

US 2004116370 PATENT INFORMATION: A1 20040617

US 2003-649106 A1 20030826 (10) APPLICATION INFO.:

> NUMBER DATE

US 2002-407375P 20020830 (60) PRIORITY INFORMATION:

US 2003-453999P 20030311 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: APPLICATION

TOWNSEND AND TOWNSEND AND CREW, LLP, TWO EMBARCADERO LEGAL REPRESENTATIVE:

CENTER, EIGHTH FLOOR, SAN FRANCISCO, CA, 94111-3834

NUMBER OF CLAIMS: 51 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 21 Drawing Page(s)

2307 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention provides compositions and methods for eliciting an immune response and compositions and methods for transfecting antigen

presenting cells.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCLM: 514/044.000 INCL

INCLS: 424/093.200; 514/150.000

NCL NCLM: 514/044.000

NCLS: 424/093.200; 514/150.000

ANSWER 8 OF 19 USPATFULL on STN

2004:139392 USPATFULL ACCESSION NUMBER:

Methods for treating and preventing infectious disease TITLE:

Krieg, Arthur M., Wellesley, MA, UNITED STATES Klinman, Dennis, Potomac, MD, UNITED STATES INVENTOR(S):

Steinberg, Alfred D., Potomac, MD, UNITED STATES

PATENT ASSIGNEE(S): University of Iowa Research Foundation, Iowa City, IA

(U.S. corporation)

The United States of America, as Represented by the Secretary, Dept. of Health & Human Services, Bethesda,

MD (U.S. corporation)

Coley Pharmaceutical Group, Inc., Wellesley, MA (U.S.

corporation)

NUMBER KIND DATE

A1 US 2004106568 20040603 PATENT INFORMATION:

20030725 (10) APPLICATION INFO.: US 2003-627331 A1

Continuation of Ser. No. US 2002-187489, filed on 2 Jul RELATED APPLN. INFO.:

2002, PENDING Division of Ser. No. US 2000-630319, filed on 31 Jul 2000, PENDING Division of Ser. No. US 1997-960774, filed on 30 Oct 1997, GRANTED, Pat. No. US

6239116 Continuation-in-part of Ser. No. US

1996-738652, filed on 30 Oct 1996, GRANTED, Pat. No. US

6207646 Continuation-in-part of Ser. No. US

1995-386063, filed on 7 Feb 1995, GRANTED, Pat. No. US

6194388 Continuation-in-part of Ser. No. US 1994-276358, filed on 15 Jul 1994, ABANDONED

DOCUMENT TYPE: Utility FILE SEGMENT: APPLICATION

> 571-272-2528 Searcher : Shears

LEGAL REPRESENTATIVE: Helen C. Lockhart, Wolf, Greenfield & Sacks, P.C.,

Federal Reserve Plaza, 600 Atlantic Avenue, Boston, MA,

02210

NUMBER OF CLAIMS: 41

EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 19 Drawing Page(s)

LINE COUNT: 3441

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Nucleic acid sequences containing unmethylated CpG dinucleotides that modulate an immune response including stimulating a Th1 pattern of immune activation, cytokine production, NK lytic activity, and B cell

proliferation are disclosed. The sequences are also useful as a

synthetic adjuvant.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCL INCLM: 514/044.000

NCL

INCLS: 536/023.100 NCLM: 514/044.000

NCLS: 536/023.100

L9 ANSWER 9 OF 19 USPATFULL on STN

ACCESSION NUMBER: 2004:120447 USPATFULL

TITLE: DNA sequences from staphylococcus aureus bacteriophage

44AHJD that encode anti-microbial polypeptides

INVENTOR(S): Pelletier, Jerry, Baie-D'Urfe, CANADA

Gros, Philippe, Lambert, CANADA Dubow, Michael, Montreal, CANADA

NUMBER DATE

PRIORITY INFORMATION: US 1999-168777P 19991201 (60)

DOCUMENT TYPE: Utility
FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: FOLEY & LARDNER, P.O. BOX 80278, SAN DIEGO, CA,

92138-0278

NUMBER OF CLAIMS: 110 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 16 Drawing Page(s)

LINE COUNT: 9802

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The disclosure concerns particular bacteriophage open reading frame, and portions and products of those open reading frames which have

antimicrobial activity Also disclosed is an S. aureus protein that interacts with an inhibitory phage protein. Methods of using such

products are also described.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCL INCLM: 435/006.000

NCL NCLM: 435/005.000 NCLS: 435/006.000 NCLS: 435/005.000

ANSWER 10 OF 19 USPATFULL on STN

2004:114692 USPATFULL ACCESSION NUMBER:

Methods of treating cancer using immunostimulatory TITLE:

oligonucleotides

INVENTOR(S): Krieg, Arthur M., Wellesley, MA, UNITED STATES

Weiner, George, Iowa City, IA, UNITED STATES

PATENT ASSIGNEE(S): University of Iowa Research Foundation, Iowa City, IA

(U.S. corporation)

DATE NUMBER KIND

US 2004087538 A1 US 2003-719493 A1 20040506 PATENT INFORMATION:

20031121 (10) APPLICATION INFO.:

Continuation of Ser. No. US 1999-337619, filed on 21 RELATED APPLN. INFO.: Jun 1999, GRANTED, Pat. No. US 6653292 Division of Ser.

No. US 1997-960774, filed on 30 Oct 1997, GRANTED, Pat. No. US 6239116 Continuation-in-part of Ser. No. US

1996-738652, filed on 30 Oct 1996, GRANTED, Pat. No. US

6207646 Continuation-in-part of Ser. No. US

1995-386063, filed on 7 Feb 1995, GRANTED, Pat. No. US

6194388 Continuation-in-part of Ser. No. US 1994-276358, filed on 15 Jul 1994, ABANDONED

DOCUMENT TYPE: Utility APPLICATION FILE SEGMENT:

Helen C. Lockhart, Ph.D., Wolf, Greenfield & Sacks, LEGAL REPRESENTATIVE:

P.C., 600 Atlantic Avenue, Boston, MA, 02210

NUMBER OF CLAIMS: 41 EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 19 Drawing Page(s)

LINE COUNT: 3433

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Nucleic acid sequences containing unmethylated CpG dinucleotides that modulate an immune response including stimulating a Th1 pattern of immune activation, cytokine production, NK lytic activity, and B cell proliferation are disclosed. The sequences are also useful as a

synthetic adjuvant.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCLM: 514/044.000 INCL

INCLS: 435/006.000; 536/023.100

NCL NCLM: 514/044.000

NCLS: 435/006.000; 536/023.100

ANSWER 11 OF 19 USPATFULL on STN

2003:312140 USPATFULL ACCESSION NUMBER:

TITLE: Novel DNA polymerase III holoenzyme delta subunit

nucleic acid molecules and proteins

Bullard, James M., Longmont, CO, UNITED STATES INVENTOR(S):

Janjic, Nebojsa, Boulder, CO, UNITED STATES McHenry, Charles S., Denver, CO, UNITED STATES

NUMBER KIND DATE _____ US 2003219737 A1 PATENT INFORMATION: 20031127 US 2001-906179 20010716 (9) APPLICATION INFO.: A1

> Shears 571-272-2528 Searcher :

RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 2001-818780, filed

on 28 Mar 2001, PENDING

NUMBER DATE

PRIORITY INFORMATION: US 2000-218246P 20000714 (60) US 2000-192736P 20000328 (60)

DOCUMENT TYPE: Utility
FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: SWANSON & BRATSCHUN L.L.C., 1745 SHEA CENTER DRIVE,

SUITE 330, HIGHLANDS RANCH, CO, 80129

NUMBER OF CLAIMS: 76 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 59 Drawing Page(s)

LINE COUNT: 14551

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Gene and amino acid sequences encoding DNA polymerase III holoenzyme δ subunits and structural genes from bacteria are provided. Also provided are antibodies and other reagents useful to identify DNA polymerase III δ subunit molecules. Also provided are methods to identify DNA polymerase III δ subunit molecules. The use of DNA polymerase III δ subunit molecules in assays to identify candidate antibiotics are also provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCL INCLM: 435/006.000

INCLS: 435/199.000; 702/020.000

NCL NCLM: 435/006.000

NCLS: 435/199.000; 702/020.000

L9 ANSWER 12 OF 19 USPATFULL on STN

ACCESSION NUMBER: 2003:309071 USPATFULL

TITLE: Method of treating cancer using immunostimulatory

oligonucleotides

INVENTOR(S): Krieg, Arthur M., Iowa City, IA, United States

Weiher, George, Iowa City, IA, United States

PATENT ASSIGNEE(S): University of Iowa Research Foundation, Iowa City, IA,

United States (U.S. corporation)

RELATED APPLN. INFO.: Division of Ser. No. US 1997-960774, filed on 30 Oct 1997, now patented, Pat. No. US 6239116, issued on 29

May 2001 Continuation-in-part of Ser. No. US

1996-738652, filed on 30 Oct 1996, now patented, Pat.

No. US 6207646, issued on 27 Mar 2001

Continuation-in-part of Ser. No. US 1995-386063, filed

on 7 Feb 1995, now patented, Pat. No. US 6194388, issued on 27 Feb 2001 Continuation-in-part of Ser. No.

US 1994-276358, filed on 15 Jul 1994, now abandoned

DOCUMENT TYPE: Utility FILE SEGMENT: GRANTED

PRIMARY EXAMINER: Martinell, James

LEGAL REPRESENTATIVE: Wolf, Greenfield & Sacks, P.C.

NUMBER OF CLAIMS: 57 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 19 Drawing Figure(s); 19 Drawing Page(s)

LINE COUNT: 3666

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Nucleic acid sequences containing unmethylated CpG dinucleotides that modulate an immune response including stimulating a Th1 pattern of immune activation, cytokine production, NK lytic activity, and B cell proliferation are disclosed. The sequences are also useful a synthetic adjuvant.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCL INCLM: 514/044.000 INCLS: 536/023.100 NCL NCLM: 514/044.000 NCLS: 536/023.100

L9 ANSWER 13 OF 19 USPATFULL on STN

ACCESSION NUMBER: 2003:271471 USPATFULL

TITLE: Methods for treating and preventing infectious disease

INVENTOR(S): Krieg, Arthur M., Wellesley, MA, UNITED STATES

Klinman, Dennis, Potomac, MD, UNITED STATES

Steinberg, Alfred D., Potomac, MD, UNITED STATES

PATENT ASSIGNEE(S): University of Iowa Research Foundation, Iowa City, IA

(U.S. corporation)

RELATED APPLN. INFO.: Continuation of Ser. No. US 2002-187489, filed on 2 Jul

2002, PENDING Division of Ser. No. US 2000-630319, filed on 31 Jul 2000, PENDING Division of Ser. No. US 1997-960774, filed on 30 Oct 1997, GRANTED, Pat. No. US

6239116 Continuation-in-part of Ser. No. US

1996-738652, filed on 30 Oct 1996, GRANTED, Pat. No. US

6207646 Continuation-in-part of Ser. No. US

1995-386063, filed on 7 Feb 1995, GRANTED, Pat. No. US

6194388 Continuation-in-part of Ser. No. US 1994-276358, filed on 15 Jul 1994, ABANDONED

DOCUMENT TYPE: Utility FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: Helen C. Lockhart, Wolf, Greenfield & Sacks, P.C.,

Federal Reserve Plaza, 600 Atlantic Avenue, Boston, MA,

02210

NUMBER OF CLAIMS: 41 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 19 Drawing Page(s)

LINE COUNT: 3449

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Nucleic acid sequences containing unmethylated CpG dinucleotides that modulate an immune response including stimulating a Th1 pattern of immune activation, cytokine production, NK lytic activity, and B cell proliferation are disclosed. The sequences are also useful as a synthetic adjuvant.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCL INCLM: 514/044.000 INCLS: 536/023.200 NCL NCLM: 514/044.000 NCLS: 536/023.200

INVENTOR(S):

L9 ANSWER 14 OF 19 USPATFULL on STN

ACCESSION NUMBER: 2003:240330 USPATFULL

TITLE: Nucleic acid and amino acid sequences relating to

Enterococcus faecalis for diagnostics and therapeutics Doucette-Stamm, Lynn A., 14 Flanagan Dr., Framingham,

MA, United States 01701

Bush, David, 205 Holland St., Somerville, MA, United

States 02144

NUMBER DATE

PRIORITY INFORMATION: US 1997-55778P 19970815 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: GRANTED

PRIMARY EXAMINER: Mosher, Mary E.

LEGAL REPRESENTATIVE: Genome Therapeutics Corporation

NUMBER OF CLAIMS: 19 EXEMPLARY CLAIM: 1,5,14

NUMBER OF DRAWINGS: 0 Drawing Figure(s); 0 Drawing Page(s)

LINE COUNT: 13738

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The invention provides isolated polypeptide and nucleic acid sequences derived from Enterococcus faecalis that are useful in diagnosis and therapy of pathological conditions; antibodies against the polypeptides; and methods for the production of the polypeptides. The invention also provides methods for the detection, prevention and treatment of pathological conditions resulting from bacterial infection.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCL INCLM: 435/320.100

INCLS: 536/023.700; 536/024.320; 435/252.300; 435/069.100; 435/006.000

NCL NCLM: 435/320.100

NCLS: 435/006.000; 435/069.100; 435/252.300; 536/023.700; 536/024.320

L9 ANSWER 15 OF 19 USPATFULL on STN

ACCESSION NUMBER: 2003:187843 USPATFULL

TITLE: DNA replication proteins of Gram positive bacteria and

their use to screen for chemical inhibitors

INVENTOR(S): O'Donnell, Michael E., Hastings-on-Hudson, NY, UNITED

STATES

Zhang, Dan, Forest Hills, NY, UNITED STATES Whipple, Richard, Elizabeth, NJ, UNITED STATES

NUMBER KIND DATE

PATENT INFORMATION: US 2003129633 A1 20030710 APPLICATION INFO.: US 2002-282287 A1 20021028 (10)

RELATED APPLN. INFO.: Continuation of Ser. No. US 1999-235245, filed on 22

Jan 1999, ABANDONED

NUMBER DATE _____

US 1998-74572P 19980213 (60) US 1998-93727P 19980722 (60) PRIORITY INFORMATION:

DOCUMENT TYPE: Utility FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: Michael L. Goldman, NIXON PEABODY LLP, Clinton Square,

P.O. Box 31051, Rochester, NY, 14603-1051

NUMBER OF CLAIMS: 32 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 11 Drawing Page(s)

LINE COUNT: 3756

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The duplex DNA of chromosomes is replicated in a multicomponent process. A helicase unwinds the DNA, a replicase synthesizes new DNA, and primase repeatedly synthesizes new primed starts on the lagging strand. The present invention is directed to the genes from Gram positive bacterium

encoding these proteins, and their characterization.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCLM: 435/006.000 TNCL

INCLS: 435/069.300; 435/199.000; 435/252.300; 435/320.100; 536/023.700

NCLM: 435/006.000 NCL

NCLS: 435/069.300; 435/199.000; 435/252.300; 435/320.100; 536/023.700

L9 ANSWER 16 OF 19 USPATFULL on STN

ACCESSION NUMBER: 2003:169096 USPATFULL

TITLE: Nucleic acid sequences and expression system relating

to Enterococcus faecium for diagnostics and

therapeutics

Doucette-Stamm, Lynn A., Framingham, MA, United States INVENTOR(S):

Bush, David, Somerville, MA, United States

Genome Therapeutics Corporation, Waltham, MA, United PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE ______ US 6583275 B1 20030624 US 1998-107532 19980630 PATENT INFORMATION:

APPLICATION INFO.: 19980630 (9)

NUMBER DATE _____

US 1998-85598P 19980514 (60) US 1997-51571P 19970702 (60) PRIORITY INFORMATION:

DOCUMENT TYPE: Utility FILE SEGMENT: GRANTED

PRIMARY EXAMINER: Marschel, Ardin H.

LEGAL REPRESENTATIVE: Genome Therapeutics Corporation

NUMBER OF CLAIMS: 34 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 0 Drawing Figure(s); 0 Drawing Page(s)

LINE COUNT: 15265

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The invention provides isolated polypeptide and nucleic acid sequences derived Enterococcus faecium that are useful in diagnosis and therapy of pathological conditions; antibodies against the polypeptides; and methods for the production of the polypeptides. The invention also provides methods for the detection, prevention and treatment of pathological conditions resulting from bacterial infection.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCLM: 536/023.100

INCLS: 435/006.000; 435/243.000; 435/320.100; 435/325.000; 536/024.300;

536/024.320

NCLM: 536/023.100 NCL

NCLS: 435/006.000; 435/243.000; 435/320.100; 435/325.000; 536/024.300;

536/024.320

ANSWER 17 OF 19 USPATFULL on STN

2003:130010 USPATFULL ACCESSION NUMBER:

Nucleic acid and amino acid sequences relating to TITLE:

Acinetobacter baumannii for diagnostics and

INVENTOR(S): Breton, Gary, Marlborough, MA, United States

Bush, David, Somerville, MA, United States

Genome Therapeutics Corporation, Waltham, MA, United PATENT ASSIGNEE(S):

States (U.S. corporation)

NUMBER KIND DATE _____ US 6562958 B1 20030513 US 1999-328352 19990604 (9) PATENT INFORMATION: APPLICATION INFO.:

NUMBER DATE _____

PRIORITY INFORMATION: US 1998-88701P 19980609 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: GRANTED
PRIMARY EXAMINER: Borin, Michael

LEGAL REPRESENTATIVE: Genome Therapeutics Corporation

NUMBER OF CLAIMS: 15 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 0 Drawing Figure(s); 0 Drawing Page(s)

16618 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The invention provides isolated polypeptide and nucleic acid sequences derived from Acinetobacter mirabilis that are useful in diagnosis and therapy of pathological conditions; antibodies against the polypeptides; and methods for the production of the polypeptides. The invention also provides methods for the detection, prevention and treatment of pathological conditions resulting from bacterial infection.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCLM: 536/023.700 INCL INCLS: 536/023.100 NCLM: 536/023.700 NCL NCLS: 536/023.100

L9 ANSWER 18 OF 19 USPATFULL on STN

ACCESSION NUMBER: 2002:343879 USPATFULL TITLE: Novel Polynucleotides

INVENTOR(S): Nakagawa, Satoshi, Tokyo, JAPAN

Mizoguchi, Hiroshi, Tokyo, JAPAN

Ando, Seiko, Tokyo, JAPAN
Hayashi, Mikiro, Tokyo, JAPAN
Ochiai, Keiko, Tokyo, JAPAN
Yokoi, Haruhiko, Tokyo, JAPAN
Tateishi, Naoko, Tokyo, JAPAN
Senoh, Akihiro, Tokyo, JAPAN
Ikeda, Masato, Tokyo, JAPAN
Ozaki, Akio, Hofu-shi, JAPAN

DOCUMENT TYPE: Utility
FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: NIXON & VANDERHYE P.C., 8th Floor, 1100 North Glebe

Road, Arlington, VA, 22201

NUMBER OF CLAIMS: 68 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 4 Drawing Page(s)

LINE COUNT: 13673

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Novel polynucleotides derived from microorganisms belonging to coryneform bacteria and fragments thereof, polypeptides encoded by the polynucleotides and fragments thereof, polynucleotide arrays comprising the polynucleotides and fragments thereof, recording media in which the nucleotide sequences of the polynucleotide and fragments thereof have been recorded which are readable in a computer, and use of them.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

INCL INCLM: 435/006.000

INCLS: 435/091.200; 435/287.200

NCL NCLM: 435/006.000

NCLS: 435/091.200; 435/287.200

L9 ANSWER 19 OF 19 USPATFULL on STN

ACCESSION NUMBER: 2002:32536 USPATFULL

TITLE: Compositions and methods for in vivo delivery of

polynucleotide-based therapeutics

INVENTOR(S): Manthorpe, Marston, San Diego, CA, UNITED STATES

Hartikka, Jukka, San Diego, CA, UNITED STATES Sukhu, Loretta, San Diego, CA, UNITED STATES

PATENT ASSIGNEE(S): Vical Incorporated, San Diego, CA (U.S. corporation)

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KIND
                            NUMBER
                                                DATE
                       ______
PATENT INFORMATION:
                       US 2002019358
                                        A1
                                               20020214
APPLICATION INFO .:
                       US 2001-839574
                                         A1
                                               20010423 (9)
                                           DATE
                             NUMBER
                       ______
                       US 2000-198823P
PRIORITY INFORMATION:
                                          20000421 (60)
                       US 2000-253153P
                                         20001128 (60)
DOCUMENT TYPE:
                       Utility
                       APPLICATION
FILE SEGMENT:
                       STERNE, KESSLER, GOLDSTEIN & FOX PLLC, 1100 NEW YORK
LEGAL REPRESENTATIVE:
                       AVENUE, N.W., SUITE 600, WASHINGTON, DC, 20005-3934
NUMBER OF CLAIMS:
                       163
EXEMPLARY CLAIM:
                       29 Drawing Page(s)
NUMBER OF DRAWINGS:
                       4605
LINE COUNT:
CAS INDEXING IS AVAILABLE FOR THIS PATENT.
      The present invention relates to pharmaceutical compositions and methods
AΒ
      to improve expression of exogenous polypeptides into vertebrate cells in
      vivo, utilizing delivery of polynucleotides encoding such polypeptides.
      More particularly, the present invention provides the use of salts, in
      particular sodium and potassium salts of phosphate, in aqueous solution,
      and auxiliary agents, in particular detergents and surfactants, in
      pharmaceutical compositions and methods useful for direct
      polynucleotide-based polypeptide delivery into the cells of vertebrates.
CAS INDEXING IS AVAILABLE FOR THIS PATENT.
TNCL
      INCLM: 514/044.000
      NCLM: 514/044.000
NCL
     (FILE 'CAPLUS, MEDLINE, BIOSIS, EMBASE, WPIDS, CONFSCI, SCISEARCH,
     JICST-EPLUS, JAPIO, USPATFULL' ENTERED AT 09:58:41 ON 04 FEB 2005)
          4314 S ("ODONNEL M"? OR "O DONNELL M"?)/AU
L10
                                                            Author (s)
           194 S "BRUCK I"?/AU
L11
         20376 S "ZHANG D"?/AU
L12
           370 S "WHIPPLE R"?/AU
L13
             2 S L10 AND L11 AND L12 AND L13
L14
L15
            71 S L10 AND (L11 OR L12 OR L13)
L16
            2 S L11 AND (L12 OR L13)
             6 S L12 AND L13
L17
            29 S (L10 OR L11 OR L12 OR L13 OR L15) AND (L1 OR L5)
L18
            31 S L14 OR L16 OR L17 OR L18
L19
            20 DUP REM L19 (11 DUPLICATES REMOVED)
L20
L20 ANSWER 1 OF 20 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 1
ACCESSION NUMBER:
                        2004:162335 CAPLUS
DOCUMENT NUMBER:
                        140:212976
                        Identification and cloning of Thermotoga maritima
TITLE:
                        deoxyribonucleate nucleotidyltransferase III \delta'
                        subunit
INVENTOR(S):
                        O'donnell, Michael E.; Yuzhakov, Alexander;
                        Yurieva, Olga; Jeruzalmi, David; Bruck, Irina
                        ; Kuriyan, John
PATENT ASSIGNEE(S):
                        USA
                        U.S. Pat. Appl. Publ., 245 pp., Cont. of U.S. Ser. No.
SOURCE:
```

716,964.

CODEN: USXXCO

DOCUMENT TYPE: Patent English LANGUAGE:

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.		DATE
US 2004038290	A1	20040226	US 2003-671419	-	20030925
US 2004043414	A1	20040304	US 2003-670844		20030925
US 2004043415	A1	20040304	US 2003-671134		20030925
US 2004106137	A1	20040603	US 2003-670817		20030925
US 2004197796	A1	20041007	US 2003-671207		20030925
US 2004048309	A1	20040311	US 2003-673098		20030926
US 2004077012	A1	20040422	US 2003-672638		20030926
US 2004081995	A1	20040429	US 2003-673127		20030926
US 2004110210	A1	20040610	US 2003-673119		20030926
PRIORITY APPLN. INFO.:			US 1997-43202P	P	19970408
			US 1998-57416	В1	19980408
			US 2000-642218	A2	20000818
			US 2000-716964	A1	20001121

The present invention relates to identification and cloning of genes AΒ encoding DNA polymerase III subunits of thermophiles. In particular, it provides identification and cloning of DNA polymerase III δ 'subunit of Thermotoga maritima for use in PCR or primer extension.

L20 ANSWER 2 OF 20 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 2

2004:162334 CAPLUS ACCESSION NUMBER:

DOCUMENT NUMBER: 140:231417

TITLE: Identification and cloning of DNA polymerase III

 δ 'subunit of Bacillus stearothermophilus

INVENTOR(S): O'donnell, Michael E.; Yuzhakov, Alexander; Yurieva, Olga; Jeruzalmi, David; Bruck, Irina

; Kuriyan, John

PATENT ASSIGNEE(S): USA

U.S. Pat. Appl. Publ., 245 pp., Cont. of U.S. Ser. No. SOURCE:

716,964.

CODEN: USXXCO

DOCUMENT TYPE: Patent English LANGUAGE:

FAMILY ACC. NUM. COUNT: 2

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	_	DATE
US 2004038289	A1	20040226	US 2003-671403		20030925
US 2004043414	A1	20040304	US 2003-670844		20030925
US 2004043415	A1	20040304	US 2003-671134		20030925
US 2004106137	A 1	20040603	US 2003-670817		20030925
US 2004197796	A 1	20041007	US 2003-671207		20030925
US 2004048309	A1	20040311	US 2003-673098		20030926
US 2004077012	A1	20040422	us 2003-672638 ·		20030926
US 2004081995	A1	20040429	US 2003-673127		20030926
US 2004110210	A1	20040610	US 2003-673119		20030926
PRIORITY APPLN. INFO.:			US 1997-43202P	P	19970408

US 1998-57416 B1 19980408 US 2000-642218 A2 20000818 US 2000-716964 A1 20001121

The present invention relates to identification and cloning of genes AB encoding DNA polymerase III subunits of thermophiles. In particular, it provides identification and cloning of DNA polymerase III δ 'subunit of Bacillus stearothermophilus for use in PCR or primer extension.

L20 ANSWER 3 OF 20 USPATFULL on STN

2004:254270 USPATFULL ACCESSION NUMBER:

Nucleic acid encoding bacillus stearothermophilus beta TITLE:

polymerase subunit

O'Donnell, Michael E., Hastings-On-Hudson, INVENTOR(S):

NY, UNITED STATES

Yuzhakov, Alexander, Malden, MA, UNITED STATES Yurieva, Olga, New York, NY, UNITED STATES Jeruzalmi, David, Cambridge, MA, UNITED STATES Bruck, Irina, New York, NY, UNITED STATES Kuriyan, John, Berkeley, CA, UNITED STATES

KIND DATE NUMBER US 2004197796 A1 20041007

PATENT INFORMATION: US 2003-671207 A1 20030925 APPLICATION INFO.:

Continuation of Ser. No. US 2000-716964, filed on 21 RELATED APPLN. INFO.: Nov 2000, PENDING Continuation-in-part of Ser. No. US

2000-642218, filed on 18 Aug 2000, PENDING Continuation

of Ser. No. US 1998-57416, filed on 8 Apr 1998,

ABANDONED

NUMBER DATE

PRIORITY INFORMATION: US 1997-43202P 19970408 (60)

DOCUMENT TYPE: ' Utility APPLICATION FILE SEGMENT:

Nixon Peabody LLP, Clinton Square, P.O. Box 31051, LEGAL REPRESENTATIVE:

Rochester, NY, 14603-1051

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 82 Drawing Page(s)

LINE COUNT: 9518

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to an isolated DNA molecule from a thermophilic bacterium which encodes a DNA polymerase III-type enzyme subunit. Also encompassed by the present invention are host cells and expression system including the heterologous DNA molecule of the present invention, as well as isolated replication enzyme subunits encoded by such DNA molecules. Also disclosed is a method of producing a recombinant thermostable DNA polymerase III-type enzyme, or subunit thereof, from a thermophilic bacterium, which is carried out by transforming a host cell with at least one heterologous DNA molecule of the present invention under conditions suitable for expression of the DNA polymerase III-type enzyme, or subunit thereof, and then isolating the DNA polymerase III-type enzyme, or subunit thereof.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L20 ANSWER 4 OF 20 USPATFULL on STN

ACCESSION NUMBER: 2004:144546 USPATFULL

Bacillus stearothermophilus SSB protein and use thereof TITLE:

O'Donnell, Michael E., Hastings-on-Hudson, INVENTOR(S):

NY, UNITED STATES

Yuzhakov, Alexander, Malden, MA, UNITED STATES Yurieva, Olga, New York, NY, UNITED STATES Jeruzalmi, David, Cambridge, MA, UNITED STATES Bruck, Irina, New York, NY, UNITED STATES Kuriyan, John, Berkeley, CA, UNITED STATES

NUMBER KIND DATE ______

PATENT INFORMATION: APPLICATION INFO.:

US 2004110210 A1 20040610 US 2003-673119 A1 20030926 (10)

Continuation of Ser. No. US 2000-716964, filed on 21 RELATED APPLN. INFO.: Nov 2000, PENDING Continuation-in-part of Ser. No. US 2000-642218, filed on 18 Aug 2000, PENDING Continuation

of Ser. No. US 1998-57416, filed on 8 Apr 1998,

ABANDONED

NUMBER DATE

PRIORITY INFORMATION:

US 1997-43202P 19970408 (60)

DOCUMENT TYPE: FILE SEGMENT:

Utility APPLICATION

LEGAL REPRESENTATIVE:

Nixon Peabody LLP, Clinton Square, P.O. Box 31051,

Rochester, NY, 14603-1051

NUMBER OF CLAIMS:

EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 82 Drawing Page(s)

LINE COUNT: 9522

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to an isolated DNA molecule from a thermophilic bacterium which encodes a DNA polymerase III-type enzyme subunit. Also encompassed by the present invention are host cells and expression system including the heterologous DNA molecule of the present invention, as well as isolated replication enzyme subunits encoded by such DNA molecules. Also disclosed is a method of producing a recombinant thermostable DNA polymerase III-type enzyme, or subunit thereof, from a thermophilic bacterium, which is carried out by transforming a host cell with at least one heterologous DNA molecule of the present invention under conditions suitable for expression of the DNA polymerase III-type enzyme, or subunit thereof, and then isolating the DNA polymerase III-type enzyme, or subunit thereof.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L20 ANSWER 5 OF 20 USPATFULL on STN

2004:138964 USPATFULL ACCESSION NUMBER:

Nucleic acid encoding bacillus stearothermophilus SSB TITLE:

protein

INVENTOR(S): O'Donnell, Michael E., Hastings-on Hudson,

NY, UNITED STATES

Yuzhakov, Alexander, Malden, MA, UNITED STATES

Yurieva, Olga, New York, NY, UNITED STATES
Jeruzalmi, David, Cambridge, MA, UNITED STATES
Bruck, Irina, New York, NY, UNITED STATES
Kuriyan, John, Berkeley, CA, UNITED STATES

RELATED APPLN. INFO.: Continuation of Ser. No. US 2000-716964, filed on 21 Nov 2000, PENDING Continuation-in-part of Ser. No. US 2000-642218, filed on 18 Aug 2000, PENDING Continuation

of Ser. No. US 1998-57416, filed on 8 Apr 1998,

ABANDONED

NUMBER DATE

PRIORITY INFORMATION: US 1997-43202P 19970408 (60)

DOCUMENT TYPE: Utility
FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: Nixon Peabody LLP, Clinton Square, P.O. Box 31051,

Rochester, NY, 14603-1051

NUMBER OF CLAIMS: 9 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 82 Drawing Page(s)

LINE COUNT: 9513

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to an isolated DNA molecule from a thermophilic bacterium which encodes a DNA polymerase III-type enzyme subunit. Also encompassed by the present invention are host cells and expression system including the heterologous DNA molecule of the present invention, as well as isolated replication enzyme subunits encoded by such DNA molecules. Also disclosed is a method of producing a recombinant thermostable DNA polyrnerase III-type enzyme, or subunit thereof, from a thermophilic bacterium, which is carried out by transforming a host cell with at least one heterologous DNA molecule of the present invention under conditions suitable for expression of the DNA polymerase III-type enzyme, or subunit thereof, and then isolating the DNA polymerase III-type enzyme, or subunit thereof.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L20 ANSWER 6 OF 20 USPATFULL on STN

ACCESSION NUMBER: 2004:107595 USPATFULL

TITLE: Bacillus stearothermophilus beta polymerase subunit and

use thereof

INVENTOR(S): O'Donnell, Michael E., Hastings-on-Hudson,

NY, UNITED STATES

Yuzhakov, Alexander, Malden, MA, UNITED STATES Yurieva, Olga, New York, NY, UNITED STATES Jeruzalmi, David, New York, NY, UNITED STATES Bruck, Irina, New York, NY, UNITED STATES Kuriyan, John, Berkeley, CA, UNITED STATES

NUMBER KIND DATE

US 2004081995 . A1 20040429 PATENT INFORMATION: US 2003-673127 A1 20030926 (10) APPLICATION INFO.:

RELATED APPLN. INFO.: Continuation of Ser. No. US 2000-716964, filed on 21

Nov 2000, PENDING Continuation-in-part of Ser. No. US 2000-642218, filed on 18 Aug 2000, PENDING Continuation

of Ser. No. US 1998-57416, filed on 8 Apr 1998,

ABANDONED

NUMBER DATE _____

US 1997-43202P 19970408 (60) PRIORITY INFORMATION:

DOCUMENT TYPE: Utility APPLICATION FILE SEGMENT:

Nixon Peabody LLP, Clinton Square, P.O. Box 31051, LEGAL REPRESENTATIVE:

Rochester, NY, 14603-1051

NUMBER OF CLAIMS: EXEMPLARY CLAIM:

82 Drawing Page(s) NUMBER OF DRAWINGS:

9515 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to an isolated DNA molecule from a thermophilic bacterium which encodes a DNA polymerase III-type enzyme subunit. Also encompassed by the present invention are host cells and expression system including the heterologous DNA molecule of the present invention, as well as isolated replication enzyme subunits encoded by such DNA molecules. Also disclosed is a method of producing a recombinant thermostable DNA polymerase III-type enzyme, or subunit thereof, from a thermophilic bacterium, which is carried out by transforming a host cell with at least one heterologous DNA molecule of the present invention under conditions suitable for expression of the DNA polymerase III-type enzyme, or subunit thereof, and then isolating the DNA polymerase III-type enzyme, or subunit thereof.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L20 ANSWER 7 OF 20 USPATFULL on STN

ACCESSION NUMBER: 2004:101150 USPATFULL

Bacillus stearothermophilus polc polymerase subunit and TITLE:

use thereof

O'Donnell, Michael E., Hastings-on-Hudson, INVENTOR(S):

NY, UNITED STATES

Yuzhakov, Alexander, Malden, MA, UNITED STATES Yurieva, Olga, New York, NY, UNITED STATES Jeruzalmi, David, Cambridge, MA, UNITED STATES Bruck, Irina, New York, NY, UNITED STATES

Kuriyan, John, Berkeley, CA, UNITED STATES

KIND DATE NUMBER _______

US 2004077012 A1 20040422 US 2003-672638 A1 20030926 (10) PATENT INFORMATION:

APPLICATION INFO.:

RELATED APPLN. INFO.: Continuation of Ser. No. US 2000-716964, filed on 21 Nov 2000, PENDING Continuation-in-part of Ser. No. US 2000-642218, filed on 18 Aug 2000, PENDING Continuation

of Ser. No. US 1998-57416, filed on 8 Apr 1998,

ABANDONED

NUMBER DATE

PRIORITY INFORMATION: US 1997-43202P 19970408 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: Nixon Peabody LLP, Clinton Square, P.O. Box 31051,

Rochester, NY, 14603-1051

NUMBER OF CLAIMS: 9 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 82 Drawing Page(s)

LINE COUNT: 9511

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to an isolated DNA molecule from a thermophilic bacterium which encodes a DNA polymerase III-type enzyme subunit. Also encompassed by the present invention are host cells and expression system including the heterologous DNA molecule of the present invention, as well as isolated replication enzyme subunits encoded by such DNA molecules. Also disclosed is a method of producing a recombinant thermostable DNA polymerase III-type enzyme, or subunit thereof, from a thermophilic bacterium, which is carried out by transforming a host cell with at least one heterologous DNA molecule of the present invention under conditions suitable for expression of the DNA polymerase III-type enzyme, or subunit thereof, and then isolating the DNA polymerase III-type enzyme, or subunit thereof.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L20 ANSWER 8 OF 20 USPATFULL on STN

ACCESSION NUMBER: 2004:63791 USPATFULL

TITLE: Thermotoga maritima delta prime polymerase subunit and

use thereof

INVENTOR(S): O'Donnell, Michael E., Hastings-on-Hudson,

NY, UNITED STATES

Yuzhakov, Alexander, Malden, MA, UNITED STATES Yurieva, Olga, New York, NY, UNITED STATES Jeruzalmi, David, Cambridge, MA, UNITED STATES Bruck, Irina, New York, NY, UNITED STATES Kuriyan, John, Riverdale, NY, UNITED STATES

NUMBER KIND DATE

PATENT INFORMATION: US 2004048309 A1 20040311 APPLICATION INFO.: US 2003-673098 A1 20030926 (10)

RELATED APPLN. INFO.: Continuation of Ser. No. US 2000-716964, filed on 21

Nov 2000, PENDING Continuation-in-part of Ser. No. US 2000-642218, filed on 18 Aug 2000, PENDING Continuation

of Ser. No. US 1998-57416, filed on 8 Apr 1998,

ABANDONED

NUMBER DATE

PRIORITY INFORMATION: US 1997-43202P 19970408 (60)

DOCUMENT TYPE: Utility
FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: Nixon Peabody LLP, Clinton Square, P.O. Box 31051,

Rochester, NY, 14603-1051

NUMBER OF CLAIMS: EXEMPLARY CLAIM: 1

82 Drawing Page(s) NUMBER OF DRAWINGS:

LINE COUNT: 8502

, CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to an isolated DNA molecule from a thermophilic bacterium which encodes a DNA polymerase III-type enzyme subunit. Also encompassed by the present invention are host cells and expression system including the heterologous DNA molecule of the present invention, as well as isolated replication enzyme subunits encoded by such DNA molecules. Also disclosed is a method of producing a recombinant thermostable DNA polymerase III-type enzyme, or subunit thereof, from a thermophilic bacterium, which is carried out by transforming a host cell with at least one heterologous DNA molecule of the present invention under conditions suitable for expression of the DNA polymerase III-type enzyme, or subunit thereof, and then isolating the DNA polymerase III-type enzyme, or subunit thereof.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L20 ANSWER 9 OF 20 USPATFULL on STN

ACCESSION NUMBER: 2004:57413 USPATFULL

Nucleic acid encoding aquifex aeolicus delta prime TITLE:

polymerase subunit

O'Donnell, Michael E., Hastings-on-Hudson, INVENTOR(S):

NY, UNITED STATES

Yuzhakov, Alexander, Malden, MA, UNITED STATES Yurieva, Olga, New York, NY, UNITED STATES Jeruzalmi, David, Cambridge, MA, UNITED STATES Bruck, Irina, New York, NY, UNITED STATES Kuriyan, John, Berkeley, CA, UNITED STATES

NUMBER KIND DATE ______ US 2004043415 A1 20040304 US 2003-671134 A1 20030925 (10)

PATENT INFORMATION: APPLICATION INFO.: RELATED APPLN. INFO.: Continuation of Ser. No. US 2000-716964, filed on 21

Nov 2000, PENDING Continuation-in-part of Ser. No. US 2000-642218, filed on 18 Aug 2000, PENDING Continuation

of Ser. No. US 1998-57416, filed on 8 Apr 1998,

ABANDONED

NUMBER DATE _____

PRIORITY INFORMATION: US 1997-43202P 19970408 (60)

DOCUMENT TYPE: Utility APPLICATION FILE SEGMENT:

LEGAL REPRESENTATIVE: Nixon Peabody LLP, Clinton Square, P.O. Box 31051,

Rochester, NY, 14603-1051

NUMBER OF CLAIMS: EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 82 Drawing Page(s)

LINE COUNT: 9517

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to an isolated DNA molecule from a AB

thermophilic bacterium which encodes a DNA polymerase III-type enzyme subunit. Also encompassed by the present invention are host cells and expression system including the heterologous DNA molecule of the present invention, as well as isolated replication enzyme subunits encoded by such DNA molecules. Also disclosed is a method of producing a recombinant thermostable DNA polymerase III-type enzyme, or subunit thereof, from a thermophilic bacterium, which is carried out by transforming a host cell with at least one heterologous DNA molecule of the present invention under conditions suitable for expression of the DNA polymerase III-type enzyme, or subunit thereof, and then isolating the DNA polymerase III-type enzyme, or subunit thereof.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L20 ANSWER 10 OF 20 USPATFULL on STN

ACCESSION NUMBER: 2004:57412 USPATFULL

TITLE: Nucleic acid encoding bacillus stearothermophilus tau

polymerase subunit

INVENTOR(S): O'Donnell, Michael E., Hastings-on- Hudson,

NY, UNITED STATES

Yuzhakov, Alexander, Malden, MA, UNITED STATES Yurieva, Olga, New York, NY, UNITED STATES Jeruzalmi, David, Cambridge, MA, UNITED STATES Bruck, Irina, New York, NY, UNITED STATES Kuriyan, John, Berkeley, CA, UNITED STATES

NUMBER KIND DATE

----US 2004043414 A1 20040304
US 2003-670844 A1 20030925 (10)

PATENT INFORMATION: US 2004043414 A1 20040304
APPLICATION INFO.: US 2003-670844 A1 20030925 (10)
RELATED APPLN. INFO.: Continuation of Ser. No. US 2000-716964, filed on 21

Nov 2000, PENDING Continuation-in-part of Ser. No. US 2000-642218, filed on 18 Aug 2000, PENDING Continuation

of Ser. No. US 1998-57416, filed on 8 Apr 1998,

ABANDONED

NUMBER DATE
----US 1997-43202P 19970408 (60)

PRIORITY INFORMATION: US 1997-4320:
DOCUMENT TYPE: Utility
FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: Nixon Peabody LLP, Clinton Square, P.O. Box 31051,

Rochester, NY, 14603-1051

NUMBER OF CLAIMS: 9 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 82 Drawing Page(s)

LINE COUNT: 9513

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to an isolated DNA molecule from a thermophilic bacterium which encodes a DNA polymerase III-type enzyme subunit. Also encompassed by the present invention are host cells and expression system including the heterologous DNA molecule of the present invention, as well as isolated replication enzyme subunits encoded by such DNA molecules. Also disclosed is a method of producing a recombinant thermostable DNA polymerase III-type enzyme, or subunit thereof, from a thermophilic bacterium, which is carried out by

transforming a host cell with at least one heterologous DNA molecule of the present invention under conditions suitable for expression of the DNA polymerase III-type enzyme, or subunit thereof, and then isolating the DNA polymerase III-type enzyme, or subunit thereof.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L20 ANSWER 11 OF 20 WPIDS COPYRIGHT 2005 THE THOMSON CORP on STN DUPLICATE 3

ACCESSION NUMBER: 2003-829557 [77] WPIDS

CROSS REFERENCE: 1999-590685 [50] C2003-233646 DOC. NO. CPI:

New DNA replication proteins (i.e. subunits of the TITLE:

> Staphylococcus aureus DNA polymerase III enzyme) and genes, useful in drug discovery to screen large libraries

of chemicals for identification of compounds with

antibiotic activity.

B04 D16 DERWENT CLASS:

O'DONNELL, M E; WHIPPLE, R; ZHANG, D INVENTOR(S):

(ODON-I) O'DONNELL M E; (WHIP-I) WHIPPLE R; (ZHAN-I) PATENT ASSIGNEE(S):

ZHANG D

COUNTRY COUNT: 1

PATENT INFORMATION:

PATENT NO KIND DATE WEEK LA PG _____ US 2003129633 A1 20030710 (200377) * 69

APPLICATION DETAILS:

PATENT NO	KIND	APPLICATION	DATE		
us 2003129633	Al Provisional Provisional Cont of	US 1998-74572P US 1998-93727P US 1999-235245 US 2002-282287	19980213 19980722 19990122 20021028		

20021028; US

PRIORITY APPLN. INFO: US 2002-282287 20021020, 1998-74572P 19980213; US 19980722; US 19990122

2003-829557 [77] WPIDS AN

1999-590685 [50] CR

US2003129633 A UPAB: 20031128 AB

NOVELTY - An isolated polypeptide, which comprises at least one functionally active subunit of a Staphylococcus aureus DNA polymerase III enzyme, is new. The subunit comprises a 573 residue dnaE amino acid sequence, a 566 residue dnaX amino acid sequence and/or a 457 residue dnaB amino acid sequence, all given in the specification.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are also included for the following:

- (1) isolated nucleic acid molecules encoding the novel subunit polypeptide or polypeptide sequence, where the nucleic acid is an isolated Staphylococcus DNA molecule encoding the dnaE, dnaX or dnaB protein or polypeptide;
 - (2) expression systems containing any of the DNA molecules of (1);

and

(3) host cells transformed with the DNA molecules of (1).

USE - The proteins and nucleic acids replicate the chromosome of Gram positive bacteria, and are useful in drug discovery to screen large libraries of chemicals for identification of compounds with antibiotic activity. Dwg.0/11

L20 ANSWER 12 OF 20 USPATFULL on STN

ACCESSION NUMBER:

2003:237815 USPATFULL

TITLE:

Methods for amplifying and sequencing nucleic acid

molecules using a three component polymerase

INVENTOR(S):

O'Donnell, Michael E., Hastings-on-Hudson,

NY, UNITED STATES

NUMBER KIND DATE _____ US 2003165972 A1 20030904 US 2003-395467 A1 20030321 (10) PATENT INFORMATION: APPLICATION INFO.:

Continuation of Ser. No. US 1999-325067, filed on 3 Jun RELATED APPLN. INFO.: 1999, GRANTED, Pat. No. US 6555349 Continuation-in-part

of Ser. No. US 1997-828323, filed on 28 Mar 1997, ABANDONED Continuation of Ser. No. US 1994-279058, filed on 22 Jul 1994, GRANTED, Pat. No. US 5668004 Continuation-in-part of Ser. No. US 1992-826926, filed

on 24 Jan 1992, ABANDONED Division of Ser. No. US

1996-696651, filed on 14 Aug 1996, ABANDONED Continuation of Ser. No. US 1994-298945, filed on 31

Aug 1994, GRANTED, Pat. No. US 5583026

DOCUMENT TYPE: Utility FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: Michael L. Goldman, NIXON PEABODY LLP, Clinton Square,

P.O. Box 31051, Rochester, NY, 14603-1051

NUMBER OF CLAIMS: 19 EXEMPLARY CLAIM: 1

INVENTOR(S):

NUMBER OF DRAWINGS: 9 Drawing Page(s)

LINE COUNT: 1544

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention is directed to a method for amplifying or sequencing a nucleic acid molecule with a three component polymerase.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L20 ANSWER 13 OF 20 USPATFULL on STN

ACCESSION NUMBER: 2003:115737 USPATFULL

TITLE: Methods for amplifying and sequencing nucleic acid

molecules using a three component polymerase O'Donnell, Michael E., Hastings-on-Hudson,

NY, United States

PATENT ASSIGNEE(S): Cornell Research Foundation, Inc., Ithaca, NY, United

States (U.S. corporation)

The Rockefeller University, New York, NY, United States

(U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION: US 6555349 B1 20030429 APPLICATION INFO.: US 1999-325067 19990603 (9)

RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 1997-828323, filed

on 28 Mar 1997 Continuation of Ser. No. US 1994-279058, filed on 22 Jul 1994, now patented, Pat. No. US 5668004 Continuation-in-part of Ser. No. US 1993-826926, filed on 22 Jan 1993, now abandoned Continuation-in-part of Ser. No. US 325067 Continuation-in-part of Ser. No. US 1999-282917, filed on 31 Mar 1999, now patented, Pat. No. US 6221642 Division of Ser. No. US 1996-696651, filed on 14 Aug 1996 Continuation of Ser. No. US 1994-298945, filed on 31 Aug 1994, now patented, Pat.

No. US 5583026

DOCUMENT TYPE: Utility

FILE SEGMENT: GRANTED

PRIMARY EXAMINER: Siew, Jeffrey LEGAL REPRESENTATIVE: Nixon Peabody LLP

NUMBER OF CLAIMS: 42 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 9 Drawing Figure(s); 9 Drawing Page(s)

LINE COUNT: 1755

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention is directed to a method for amplifying or sequencing a nucleic acid molecule with a three component polymerase comprising a DNA polymerase component, a sliding clamp component, and a

clamp loader component.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L20 ANSWER 14 OF 20 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 4

ACCESSION NUMBER: 2002:371330 CAPLUS

DOCUMENT NUMBER: 137:274946

TITLE: Analysis of a multicomponent thermostable DNA

polymerase III replicase from an extreme thermophile

AUTHOR(S): Bruck, Irina; Yuzhakov, Alexander; Yurieva,

Olga; Jeruzalmi, David; Skangalis, Maija; Kuriyan,

John; O'Donnell, Mike

CORPORATE SOURCE: Howard Hughes Medical Institute, New York, NY, 10021,

USA

SOURCE: Journal of Biological Chemistry (2002), 277(19),

17334-17348

CODEN: JBCHA3; ISSN: 0021-9258

PUBLISHER: American Society for Biochemistry and Molecular

Biology

DOCUMENT TYPE: Journal LANGUAGE: English

AB This report takes a proteomic/genomic approach to characterize the DNA polymerase III replication apparatus of the extreme thermophile, Aquifex aeolicus. Genes (dnaX, holA, and holB) encoding the subunits required for

clamp loading activity $(\tau, \delta, \text{ and } \delta')$ were identified.

The dnaX gene produces only the full-length product, τ , and therefore differs from Escherichia coli dnaX that produces two proteins (γ and

 $\tau)$. Nonetheless, the A. aeolicus proteins form a $\tau\delta\delta$

complex. The dnaN gene encoding the β clamp was identified, and the $\tau\delta\delta^{\,\prime}$ complex is active in loading

 β onto DNA. A. aeolicus contains one dnaE homolog, encoding the

 α subunit of DNA polymerase III. Like E. coli, A. aeolicus α and τ interact, although the interaction is not as tight as the α - τ contact in E. coli. In addition, the A. aeolicus homolog to dnaQ, encoding the & proofreading 3'-5'-exonuclease, interacts with α but does not form a stable $\alpha \cdot \epsilon$ complex, suggesting a need for a brace or bridging protein to tightly couple the polymerase and exonuclease in this system. Despite these differences to the E. coli system, the A. aeolicus proteins function to yield a robust replicase that retains significant activity at 90°. Similarities and differences between the A. aeolicus and E. coli pol III systems are discussed, as is application of thermostable pol III to biotechnol. THERE ARE 60 CITED REFERENCES AVAILABLE FOR THIS

REFERENCE COUNT: RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L20 ANSWER 15 OF 20 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 5

ACCESSION NUMBER:

2001:101168 CAPLUS

DOCUMENT NUMBER:

134:143863

TITLE:

DNA replication proteins of Gram-positive bacteria and

their use to screen for chemical inhibitors

INVENTOR(S):

O'donnell, Michael E.; Bruck, Irina

; Zhang, Dan; Whipple, Richard The Rockefeller University, USA

PATENT ASSIGNEE(S): SOURCE:

PCT Int. Appl., 238 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO.	KIND DATE	Al	PPLICATION NO.	DATE				
WO 2001009164	A2 2001	0208 W	2000-US20666	20000728				
• • • •		•		CH, CN, CR, CU,				
CZ, DE, DK,	DM, EE, ES,	FI, GB, G	GD, GE, GH, GM,	HR, HU, ID, IL,				
IN, IS, JP,	KE, KG, KP,	KR, KZ, 1	LC, LK, LR, LS,	LT, LU, LV, MA,				
MD, MG, MK,	MN, MW, MX,	NO, NZ,	PL, PT, RO, RU,	SD, SE, SG, SI,				
SK, SL, TJ								
RW: GH, GM, KE,	LS, MW, MZ,	SD, SL, S	SZ, TZ, UG, ZW,	AT, BE, CH, CY,				
DE, DK, ES,	FI, FR, GB,	GR, IE,	IT, LU, MC, NL,	PT, SE, BF, BJ,				
CF, CG, CI,	CM, GA, GN,	GW, ML, N	MR, NE, SN, TD,	TG				
AU 2000067499	A5 2001	0219 A	J 2000-67499	20000728				
US 2003129633	A1 2003	A1 20030710 US 2002-282287						
PRIORITY APPLN. INFO.:		U:	A 19990122					
	•	U:	S 1999-146178P	P 19990729				
		U:	S 1998-74522P	P 19980127				
		U:	S 1998-93727P	P 19980722				
		W	WO 2000-US20666 W 2000072					

ΑB The present invention relates to α -large, α -small, δ , δ' , τ , β , SSB, DnaG, and DnaB and (polC, dnaE, holA, holB, dnaX, dnaN, ssb, dnaG, dnaB) genes encoding them from Gram pos. bacteria, preferably Streptococcus pyogenes and Staphylococcus aureus. The individual genes and proteins or polypeptides are useful in identification of compds. with antibiotic activity. Thus, the structure and mechanism of the chromosomal replicase of S. pyogenes and S. aureus have been elucidated. These DNA polymerases use a sliding clamp (the dnaN-encoded β subunit) and clamp loader (the dnaX-encoded

> 571-272-2528 Searcher : Shears

 τ subunit). The clamp and clamp loader components of Gram-neg. cells could be exchanged for those of Gram-pos. cells.

L20 ANSWER 16 OF 20 USPATFULL on STN

ACCESSION NUMBER:

2001:59664 USPATFULL

TITLE:

Process for reconstituting the polymerase III* and other subassemblies of E. coli DNA polymerase III

holoenzyme from peptide subunits

INVENTOR(S):

O'Donnell, Michael E., Hastings-on-Hudson,

NY, United States

PATENT ASSIGNEE(S):

Cornell Research Foundation, Inc., Ithaca, NY, United

States (U.S. corporation)

NUMBER KIND DATE

PATENT INFORMATION:

US 6221642 B1 20010424 US 1999-282917 19990331 (9)

APPLICATION INFO.: RELATED APPLN. INFO.:

Division of Ser. No. US 1996-696651, filed on 14 Aug

1996, now abandoned Continuation of Ser. No. US

1994-298945, filed on 31 Aug 1994, now patented, Pat.

No. US 5583026

DOCUMENT TYPE:

Utility Granted

FILE SEGMENT:

Granted

PRIMARY EXAMINER: LEGAL REPRESENTATIVE: Weber, Jon P. Nixon Peabody LLP

NUMBER OF CLAIMS:

7

EXEMPLARY CLAIM:

1
29 Drawing Figure(s); 14 Drawing Page(s)

NUMBER OF DRAWINGS: LINE COUNT:

1890

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AΒ

The process of the invention provides for the reconstitution of the polymerase III* subassembly, Pol III*, of E. coli DNA polymerase III holoenzyme from substantially pure peptide subunits. In the first of two general schemes in which the subunits are added in a specified order,

 γ and τ are premixed before addition of δ and δ' .

In the second general scheme, δ ' is first assembled onto γ (or τ); then the excess δ ' is removed before adding τ (or γ), following which δ is added. Reconstituted Pol III* had

the same subunit composition as purified natural Pol III*, as well as similar activity. Other smaller subassemblies of the polymerase III holoenzyme may also be reconstituted by the process of the invention.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L20 ANSWER 17 OF 20 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 6

ACCESSION NUMBER:

1999:673058 CAPLUS

DOCUMENT NUMBER:

INVENTOR(S):

131:319658

TITLE:

DNA polymerase holoenzyme III derived from

thermophilic organisms that functions as a chromosomal

replicase, and the genes encoding its subunits Yurieva, Olga; Kuriyan, John; O'Donnell, Michael

E.; Jeruzalmi, David

PATENT ASSIGNEE(S):

The Rockefeller University, USA

SOURCE:

PCT Int. Appl., 156 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

KIND DATE APPLICATION NO. DATE PATENT NO. _____ ____ -----_____ -----WO 1998-US7070 A1 19991021 19980409 WO 9953074 W: AL, AU, BA, BB, BG, BR, CA, CN, CU, CZ, EE, GE, GW, HU, ID, IL, IS, JP, KP, KR, LC, LK, LR, LT, LV, MG, MK, MN, MX, NO, NZ, PL, RO, SG, SI, SK, SL, TR, TT, UA, UZ, VN, YU, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM RW: GH, GM, KE, LS, MW, SD, SZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG 19991101 AU 1998-71057 AU 9871057 **A**1 19980409 A 19980409 PRIORITY APPLN. INFO.: WO 1998-US7070 A DNA Polymerase has been identified in a thermophile that functions as a chromosomal replicase. The specific enzyme is a holoenzyme III that has

AB A DNA Polymerase has been identified in a thermophile that functions as a chromosomal replicase. The specific enzyme is a holoenzyme III that has been identified in Thermus thermophilus, and corresponds to Polymerase III in Escherichia coli. The genes and the polypeptides corresponding to T. thermophilus γ , τ , ε , α , and β subunits that

they encode are disclosed, as are probes, vectors, methods of preparation

and

the methods of use. The enzymes of the present invention and their components are particularly well suited for use in procedures for the preparation of DNA, such as PCR, because of the speed and accuracy that they are able to achieve.

REFERENCE COUNT:

THERE ARE 1 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L20 ANSWER 18 OF 20 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 7

ACCESSION NUMBER:

1999:487306 CAPLUS

DOCUMENT NUMBER:

131:113137

TITLE:

DNA replication proteins of gram positive bacteria and

their use to screen for chemical inhibitors

INVENTOR(S):

O'Donnell, Michael E.; Zhang, Dan;

Whipple, Richard

PATENT ASSIGNEE(S):

The Rockefeller University, USA

SOURCE:

PCT Int. Appl., 132 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent English

LANGUAGE:

English

1

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO.				KIND DATE		APPLICATION NO.						DATE					
WO 9937661			A1 19990729			WO 1999-US1547					19990125						
		ΑU,	•	•													
	RW:	AT,	BE,	CH,	CY,	DE,	DK,	ES,	FI,	FR,	GB,	GR,	ΙE,	IT,	LU,	MC,	NL,
		PT,	SE														
CA	2318	574			AΑ		1999	0729		CA 1	999-	2318	574		1:	9990	125
ΑU	9923	416			A1		1999	0809		AU 1	999-	2341	6		1:	9990	125
ΕP	1056	763			A1		2000	1206		EP 1	999-	9033	77		1	9990	125
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		IE,	FI														

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    US 2003129633
                       A1
                               20030710
                                          US 2002-282287
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PRIORITY APPLN. INFO.:
                                          US 1998-74522P
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                                                            B1 19990122
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                                          WO 1999-US1547
                                                            W 19990125
    The duplex DNA of chromosomes is replicated in a multicomponent process.
AΒ
    A helicase unwinds the DNA, a replicase synthesizes new DNA, and primase
     repeatedly synthesizes new primed starts on the lagging strand. The
    present invention is directed to the genes from gram-pos. bacterium
     encoding these proteins, and their characterization. Gene sequences are
    provided for the dnaE (\alpha-subunit), dnaX (gamma/tau subunit), dnaB
     (helicase), polC (DNA polymerase III-L), dnaN (β-subunit),
     and dnaG (primase) genes from Staphylococcus aureus. The invention dets.
     that the replicase of Staphylococcus operates as a 3-component system in
    which a clamp loader enzyme assembles a sliding clamp protein onto DNA.
    The sliding clamp then binds the DNA polymerase III holoenzyme making it
    highly efficient. The invention identifies two DNA polymerase III enzymes
     in gram-pos. bacterium, each of which operate with the clamp and clamp
     loader, to extend a single primed site around a long (>5 kb)
     single-stranded DNA template. These replication proteins can be utilized
     in a variety of assays to screen chemical compound libraries for an
antibiotic
     compound
                              THERE ARE 5 CITED REFERENCES AVAILABLE FOR THIS
REFERENCE COUNT:
                        5
                              RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT
L20 ANSWER 19 OF 20 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 8
                        1998:682540 CAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                        129:299056
TITLE:
                        A DNA polymerase III homolog of the thermophilic
                        bacterium Thermus thermophilus involved in chromosomal
                        replication
                        Yurieva, Olga; Kuriyan, John; O'donnell, Michael
INVENTOR(S):
                        E.; Jeruzalmi, David
                        The Rockefeller University, USA
PATENT ASSIGNEE(S):
                        PCT Int. Appl., 154 pp.
SOURCE:
                        CODEN: PIXXD2
DOCUMENT TYPE:
                        Patent
                        English
LANGUAGE:
FAMILY ACC. NUM. COUNT: 1
PATENT INFORMATION:
                   KIND DATE APPLICATION NO.
                                                               DATE
    PATENT NO.
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    WO 9845452 A2 19981015 WO 1998-US6921
WO 9845452 A3 19981217
                                                                 19980408
        W: AL, AU, BA, BB, BG, BR, CA, CN, CU, CZ, EE, GE, GW, HU, ID, IL,
            IS, JP, KP, KR, LC, LK, LR, LT, LV, MG, MK, MN, MX, NO, NZ, PL,
            RO, SG, SI, SK, SL, TR, TT, UA, UZ, VN, YU, AM, AZ, BY, KG, KZ,
            MD, RU, TJ, TM
        RW: GH, GM, KE, LS, MW, SD, SZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES,
            FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI,
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Searcher : Shears 571-272-2528

R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,

AU 1998-76839

EP 1998-924742

19980408

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CM, GA, GN, ML, MR, NE, SN, TD, TG

19981030

20000308

A1

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AU 9876839

EP 983365

IE, FI

PRIORITY APPLN. INFO.: US 1997-823407 A 19970408 WO 1998-US6921

A DNA polymerase holoenzyme III identified in Thermus thermophilus and AB that corresponds to DNA polymerase III of Escherichia coli is described for use in primer-mediated amplification of DNA. In particular, the clamp structure of DNA polymerase III can be used to extend a primer over a long stretch of single-stranded DNA. The enzyme may be obtained from a range of known thermophilic microorganisms. Genes for five subunits $(\alpha,$ $\beta,~\gamma,~\epsilon,~and~\tau)$ of the T. thermophilus holoenzyme are cloned and expressed. The enzyme is particularly well suited for use

in procedures for the preparation of DNA, such as PCR, because of the speed

and

accuracy that they are able to achieve. The dnaX gene of T. thermophilus was cloned using PCR with primers derived from conserved sequences of other known dnaX genes to generate a probe to screen an XbaI bank. The γ and τ subunits encoded by the dnaX gene were manufactured in Escherichia coli using the the pET expression system and shown to have an ATPase activity. The two subunits were synthesized from a single gene by efficient ribosomal frameshifting. Conserved sequence-derived primers were used to clone fragments of the other genes for subunits of the enzyme and these were extended to obtain full-length sequences by standard methods.

L20 ANSWER 20 OF 20 USPATFULL on STN

ACCESSION NUMBER: 96:113822 USPATFULL

Process for reconstituting the polymerase III* and TITLE:

other subassemblies of E. coli DNA polymerase III

holoenzyme from peptide subunits

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NY, United States

PATENT ASSIGNEE(S): Cornell Research Foundation, Inc., Ithaca, NY, United

States (U.S. corporation)

NUMBER KIND DATE PATENT INFORMATION: US 5583026 19961210 APPLICATION INFO.: US 1994-298945 19940831 (8) DOCUMENT TYPE: Utility DOCUMENT TYPE: FILE SEGMENT: Granted
PRIMARY EXAMINER: Wityshyn, Michael G.
ASSISTANT EXAMINER: Weber, Jon P.

LEGAL REPRESENTATIVE: Nixon, Hargrave, Devans & Doyle

NUMBER OF CLAIMS: 53 EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 29 Drawing Figure(s); 12 Drawing Page(s)

2275 LINE COUNT:

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The process of the invention provides for the reconstitution of the AB polymerase III* subassembly, Pol III*, of E. coli DNA polymerase III holoenzyme from substantially pure peptide subunits. In the first of two general schemes in which the subunits are added in a specified order, γ and τ are premixed before addition of δ and δ' . In the second general scheme, $\delta^{\,\prime}$ is first assembled onto γ (or τ); then the excess δ' is removed before adding τ (or γ), following which δ is added. Reconstituted Pol III* had the same subunit composition as purified natural Pol III*, as well as

similar activity. Other smaller subassemblies of the polymerase III holoenzyme may also be reconstituted by the process of the invention.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

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